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14. ABSTRACT The aggressive behavior of malignant breast cancer is determined by a complex array of signaling pathways that regulate cell growth, survival and migration. The PI 3-K-Akt pathway has been linked to all of these responses. The current paradigm states that deregulated PI 3-K-Akt signaling promotes cancer progression. Many of the enzymes that regulate PI 3-K signaling are frequently mutated in human breast cancer, thereby up-regulating Akt activity and increasing tumor cell growth and survival. This is best illustrated by the finding that the catalytic subunit of PI 3-K, <i>PIK3CA</i> , is the most frequently mutated oncogene in breast cancer. However, recent studies have demonstrated that distinct Akt isoforms can either inhibit or enhance breast cancer cell invasive migration and metastasis <i>in vitro</i> and <i>in vivo</i> , probably by phosphorylating a different set of substrates in an isoform specific manner. Afadin is a Ras target gene that regulates cell-cell adhesion downstream of Ras. Afadin protein loss of expression is associated with poor outcome in breast cancer patients. The most significant finding of this research thus far is that Afadin –isoform 3, which is the longer, ubiquitously expressed form of the protein, is a substrate of Akt, down stream of the PI-3K pathway.					
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Introduction:

Increased Akt activity is detected in aggressive human breast cancers [1, 2] and is associated with poor prognosis and higher probability of relapse accompanied by distant metastases in patients [3]. Experiments in transgenic mice have revealed that Akt promotes mammary tumor progression by increasing cell survival [4]. Moreover, many of the proteins in the Akt pathway are either oncogenes (e.g. Akt itself, PI 3-K, MDM2) or tumor suppressors (e.g. the PIP3 phosphatase PTEN and Akt phosphatase PH domain Leucine-rich repeat Protein Phosphatase (PHLPP)), and indeed these proteins are frequently deregulated in breast cancer. The significance of the Akt pathway in cancer has made it an attractive target for the development of small molecule inhibitors [5]. There are three mammalian Akt isoforms: Akt1 (PKB α), Akt2 (PKB β) and Akt3 (PKB γ). The three Akt isoforms are encoded by distinct genes, but share a high degree of amino acid similarity and are activated by similar mechanisms and recognize and phosphorylate the same consensus motif [6]. The objectives of this research are to discover novel substrates of the PI-3K-Akt pathway that promote malignant phenotype in breast cancer, with a special scope on Akt-isoform specific substrates and Afadin, as a putative novel substrate of PI-3K pathway.

In the previous reported I was able to show that Afadin isoform 3 is a novel substrate of all 3 Akt isoforms. The phosphorylation site is S1718, and it is sensitive to a line of PI-3K and Akt inhibitors. I also showed that following phosphorylation of Afadin it translocates into the nucleus.

Body:

In the previous years I found that Afadin isoform 3 is a novel substrate of the PI-3K pathway. I found that it is a substrate of Akt. The phosphorylation site is S1718. I was trying to find out whether it's an Akt isoform specific substrate, in this part I found that it's a general substrate that can be phosphorylated by all three Akt isoforms but not by other AGC kinases.

Next, I was focusing on trying to establish the biological significance of this phosphorylation, and to reveal the connection between PI-3K and Akt to this important adhesion protein.

In the previous years I also started the proteomic screen. I had silenced the expression of a specific kinase by the pLKO-Tet-On system. The main goal of this part of the research is to find novel isoform specific substrates of Akt that would help explain the importance of this family of kinases in many processes in breast cancer.

Description of the research accomplishment associated with the tasks outlined in the approved statement of work:

Aim1: Afadin Regulates Breast Cancer Cell Migration Downstream of Akt and SGK.

In the reported period I was focusing on writing and publishing a manuscript that summarizes my studies. The manuscript entitled "The Adherens Junction Protein Afadin is an Akt substrate that Regulates Breast Cancer Cell Migration" was published in Molecular cancer Research in November 2013 and is in the appendix.

We were able to show that Afadin is phosphorylated by Akt downstream of the PI3K at Ser1718 (Figures 1-2). This phosphorylation leads to translocation of this otherwise membranal protein into the nucleus (figures 3-5). It also leads to enhanced migration of breast cancer cells (figure 6). And Finally we were able to show that this phenomenon is relevant to real breast tumors, as it was recapitulated in tumor microarray that consists of many tumor specimens (Figure 7).

All the information and the precise methodology can be found in the manuscript.

Aim 2: Phospho-proteomic analysis of Akt1/2/3 and SGK1/2/3 substrates in breast cancer

The work on this aim was postponed in the reported year, as initial results were not convincing enough and we were not sure that the methodology that was used can produce high quality data. For this reason we decided to focus on aim 1 and publishing the manuscript.

Key Research Accomplishments:

- Afadin isoform 3 was found to be a novel substrate of Akt in the PI-3K signaling Pathway, that can be phosphorylated in S1718 by all three Akt isoforms.
- The phosphorylation of S1718 leads to a change in intracellular localization of this adhesion protein.
- Following phosphorylation the protein translocates into the nucleus, whereas the nonphosphorylatable protein is only membranous.
- Phosphorylation of Afadin isoform 3 in S1718 leads to enhanced migration of the cells.
- nuclear localization of Afadin, regulated by phosphorylation at Ser1718 by the Akt pathway, is clinically relevant for breast cancer progression, and the nuclear translocation is also seen in a subset of breast tumors.

Reportable Outcomes:

My training program in the last year included weekly meetings with Dr. Toker, my postdoctoral mentor, where we analyzed results and plan experiments. In addition, our laboratory meets once a week for a lab meeting. In these meetings we receive feedback on our work, as well as advice and input. In the last year I presented my project 6 times in lab meetings. Moreover, the Toker laboratory has a bi-weekly journal club meeting where discussions focus on recently published literature in the breast cancer field. I presented twice in the journal club and participated in all of the meetings we had. Finally, as part of the various intramural training activities within the laboratory, our lab attends a weekly cancer signaling meeting at Harvard Medical School. These meetings are designated to refine students' and postdocs' presentation skills. These meetings allow the participants and speakers to receive feedback on work from leaders in their respective fields, including Lewis Cantley, Joan Brugge and John Blenis. In the reported year I presented once in this forum.

Conclusions:

In the reported year I was finishing my experiments, repeating and refining experiments in order to publish my manuscript.

The manuscript entitled “The Adherens Junction Protein Afadin is an Akt substrate that Regulates Breast Cancer Cell Migration” was recently published in Molecular cancer research. This paper identifies Ser1718 in Afadin as a phosphorylation site of Akt downstream of the PI3K pathway. This is a novel substrate of the PI-3K / Akt pathway. I found that following the phosphorylation of S1718 there is a translocation of this big otherwise membranal protein to the nucleus. This is an interesting phenomenon because Afadin is a very big protein, that up until now was thought to reside in the membrane alone. This process of nuclear translocation could have several implications. First, it's possible that the nuclear translocation of Afadin leads to activation of transcription factor or other pathways in the nucleus. Another option is that it's the loss of the membranal signal itself that leads to the “loss of activity” of Afadin as an adhesion protein in the membrane. In addition, phosphorylation of Afadin isoform 3 in S1718 changes the migratory characteristics of cells. The non-phosphorylatable protein is less migratory than the wild type and the phosphomimetic protein. In other words, in order for the protein to serve as an adhesion protein and “protect” cells from migration, it cannot be phosphorylated. Once the protein is phosphorylated one of two things can happen, either the nuclear translocation itself leads to elevation in migration through activation of other pathways or proteins, or the loss of the membranal signal is responsible for the elevated migration of these cells. We discovered that these two scenarios co-exist and that migration is elevated by the translocation to the nucleus, and at the same time, loss of Afadin (which includes loss of the membrane and nuclear fractions) also leads to changes in cell-to-cell adhesion. In addition, we were able to show that this phenomenon is relevant to breast cancer tumors and also happens in the tumors.

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The Adherens Junction Protein Afadin is an Akt substrate that Regulates Breast Cancer Cell Migration

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23 **Abstract**

24 The PI 3-K and Akt signaling pathway regulates all phenotypes that contribute to
25 progression of human cancers, including breast cancer. Akt mediates signal relay by
26 phosphorylating numerous substrates, which are causally implicated in responses such as
27 cell growth, survival, metabolic reprogramming and migration and invasion. Here we identify a
28 new Akt substrate, the adherens junction protein Afadin, that is phosphorylated by Akt at
29 Ser1718. We show that under conditions of physiological IGF-1 signaling and oncogenic PI
30 3-K and Akt, Afadin is phosphorylated by all Akt isoforms, and that this phosphorylation
31 elicits a relocalization of Afadin from adherens junctions to the nucleus. Phosphorylation of
32 Afadin also results in a marked increase in breast cancer cell migration that is dependent on
33 Ser1718 phosphorylation. We also observe nuclear localization in breast cancer tissues,
34 indicating that regulation of Afadin by the PI 3-K and Akt pathway has pathophysiological
35 significance.

36 Implications: Phosphorylation of the adhesion protein Afadin by Akt downstream of the PI 3-
37 K pathway, leads to re-distribution of Afadin and controls cancer cell migration.

38

39 Introduction

40 The phosphoinositide 3-kinase (PI 3-K) and Akt signaling pathway orchestrates virtually
 41 all aspects of epithelial and tumor cell behavior, from initial transformation to dysplasia and
 42 ultimately the dissemination of cancer cells to distant metastatic sites (1). In addition,
 43 mutations in genes that encode proteins that are rate-limiting for transducing the PI 3-K and
 44 Akt signaling are frequent mutated in human cancers. This is most evident in breast cancer,
 45 whereby according to molecular subtype, the most frequent genetic lesions are oncogenic
 46 mutations in the p110 α PI 3-K catalytic subunit, *PIK3CA*, inactivation or loss of
 47 heterozygosity of the tumor suppressors *PTEN* and *INPP4B* and amplification or somatic
 48 activating mutations in one of the three Akt genes *AKT1*, *AKT2* and *AKT3* (2,3). All of these
 49 lesions ultimately result in hyperactivation of Akt and phosphorylation of downstream
 50 substrates that transduce the signal to secondary effector pathways and in turn the
 51 modulation of phenotypes associated with malignancy, including cell growth, proliferation,
 52 survival, metabolic reprogramming, and cell migration and invasion (4). Moreover, since
 53 most of the proteins that function to transduce PI 3-K and Akt signaling are enzymes with
 54 catalytic pockets, this pathway is highly druggable and numerous phase I and II clinical trials
 55 are underway with small molecule inhibitors targeting PI 3-K or Akt isoforms for single agent
 56 or combination therapy, including in breast cancer (5).

57 Increased Akt activity is detected in aggressive human breast cancers and is associated
 58 with poor prognosis and higher probability of relapse accompanied by distant metastases in
 59 patients (6-8). The ability of cancer cells to migrate requires signals which lead to the
 60 rearrangement of the actin cytoskeleton as well as proteolysis of the extracellular matrix
 61 (9,10). Importantly, molecular genetic as well as *in vivo* studies have demonstrated that Akt
 62 isoforms play unique roles in modulating breast cancer cell invasion leading to metastatic
 63 dissemination, such that Akt2 is a metastasis enhancer, whereas Akt1 either does not

promote metastasis or can actually block this process and thus function as a suppressor (11,12). Yet in other cell types and tissues, Akt isoforms either have no specificity in modulating cell migration, or even have opposing roles to those identified in epithelial cells, such as that reported for fibroblast migration (9). Regardless, the signaling specificity attributed to Akt isoforms highlights the importance of a complete understanding of the mechanism that govern cancer cell phenotypes such as invasive migration and metastasis, if specific drugs are to be developed for effective cancer therapy. In terms of mechanisms that explain the function of Akt in the control of migration, invasion and metastasis, a number of specific substrates have been identified recently. These include the actin-bundling protein palladin, a unique Akt1 substrate that functions to mediate the inhibitor activity of this Akt isoform in cell migration (13). Other substrates include girdin, that following phosphorylation accumulates in the leading edges of migrating cells and is essential for the integrity of the actin cytoskeleton and cell migration (9). Also included in this list are ACAP1, whose phosphorylation controls the recycling of integrin- β 1 and cell migration, and the G-protein coupled receptor EDG-1 that is required for endothelial cell chemotaxis (14,15). Recent global phospho-proteomic studies from cancer cell lines and tissues have identified thousands of novel phosphoproteins with phosphorylation sites that conform to the optimal Akt consensus motif, RxRxxS/T, greatly accelerating the discovery of Akt targets that transduce the signal (16).

Afadin, a tumor suppressor-like protein encoded by the *MLLT4* gene, is a multi-domain F-actin-binding protein that is expressed in epithelial cells, neurons, fibroblasts and endothelial cells (17,18). There exist two splice variants: l-Afadin and s-Afadin (18). The longer splice variant, l-Afadin (herein referred to as Afadin unless otherwise specified) has two Ras associating domains, a Forkhead associating domain, a Dilute domain, a PDZ domain, three proline-rich domains and the F-actin binding domain at the carboxyl-terminus

(see Fig. 1A). s-Afadin, the shorter splice variant, lacks the F-actin binding domain and the third proline-rich domain and its expression is restricted to neuronal tissues (19). Human s-Afadin is identical to the gene product of *AF6*, an ALL-1 fusion partner involved in acute myeloid leukemia (20,21).

Afadin is localized at epithelial adherens junctions (18), consisting of two adhesion complexes, the Nectin-Afadin and the E-cadherin-Catenin complexes (20). The role of Afadin in the adherens junction complex is not completely understood. Afadin interacts with cell adhesion molecules, cytoskeletal components, signaling molecules and is generally considered to function as an adaptor protein. Knockout of Afadin in mice results in embryonic lethality due to disorganization of the ectoderm, impaired migration of the mesoderm and impaired gastrulation. Moreover, loss of cell polarity due to improperly assembled adherens junction and tight junctions is observed (17,19,20,22). Afadin has also been shown to regulate integrin-mediated cell adhesion and cell migration, although it appears that the function of Afadin in positively or negatively regulating cell motility is context-dependent (23-27).

Here, we show that Afadin is a substrate of Akt whose phosphorylation leads to its relocalization from the plasma membrane to the nucleus, concomitant with an enhancement of breast epithelial and cancer cell migration.

107 **Materials and Methods**

108 **Cell culture**

109 HEK293T, HeLa, MCF7, MDA-MB-231, MDA-MB-453, MDA-MB-468, SKBR3, BT-549, ZR-
 110 75-1, MCF10A, T47D and ZR-75-30 were obtained from the American Type Culture
 111 Collection (ATCC), and authenticated using Short Tandem Repeat (STR) profiling. Cells
 112 were maintained in culture not more than 6 months. Cells were routinely screened for
 113 mycoplasma contamination. Cell lines were maintained as follows: HEK293T, HeLa, MCF7,
 114 MDA-MB-231, MDA-MB-453 and MDA-MB-468, Dulbecco's modified Eagle's medium
 115 (DMEM) (Cellgro; Manassas, VA) supplemented with 10% fetal bovine serum (FBS)
 116 (HyClone; Waltham, MA); SKBR3, McCoy's 5A medium (Cambrex; East Rutherford, NJ)
 117 supplemented with 10% FBS; BT-549 and ZR-75-1, RPMI 1640 (Cellgro) supplemented
 118 with 10% FBS; MCF10A DMEM/Ham's F12 supplemented with 5% equine serum (GIBCO;
 119 Carlsbad, CA), 10 µg/ml Insulin (Sigma-Aldrich; St. Louis, MO) 500 ng/ml hydrocortisone
 120 (Sigma-Aldrich), 20 ng/ml EGF (R&D Systems; Minneapolis, MN), and 100 ng/ml cholera
 121 toxin (List Biological Labs; Campbell, CA); T47D and ZR-75-30 in RPMI 1640
 122 supplemented with 10% FBS and 10 µg/ml Insulin.

123

124 **Growth Factors and Inhibitors**

125 Cells were stimulated with recombinant human IGF-1 (R&D Systems) at a final
 126 concentration of 100 ng/ml for 20 min unless otherwise specified. CGK733 (Sigma-Aldrich)
 127 was added to cells for 4 hr in a final concentration of 10 µM prior to IGF1 stimulation. All
 128 other inhibitors were added to cells for 20 min prior to stimulation at the following final
 129 concentrations: wortmannin (Sigma-Aldrich), 100 nM; BEZ235 (Cayman Chemical
 130 company; Ann Harbor, MI) and MK2206 (Active Biochem; Maplewood, NJ), 1 µM; A66
 131 (Symansis; Temecula, CA), 0.7 µM. Rapamycin (Sigma-Aldrich), 100 nM; PF4708671

(Sigma-Aldrich), 10 μ M. GSK650394 (R&D Systems), 5 μ M; Cycloheximide (Sigma-Aldrich) was used for 3-6 hr at 20 μ g/ml.

Antibodies

Anti-phospho-Afadin Ser1718 antibody, anti-pan-Akt antibody, anti-Akt1, anti-Akt2, anti-Akt3, anti-phospho-Akt Ser473, anti-GSK3 β , anti-phospho-GSK3 β Ser9, anti-p110 α , anti-S6K, anti-phospho-S6K Thr389, anti E-cadherin, anti-NDRG1, anti-phospho-NDRG1 Thr346, anti-CENP-A, anti-NuP98, anti-Fibrillarin and anti-Histone H3 were from Cell Signaling Technology (Danvers, MA). Anti-Myc antibody, anti-Tubulin and anti-lamin A/C antibodies were from Santa Cruz Biotechnology (Santa Cruz, CA). Anti-Afadin antibodies used for immunoblotting were from Bethyl laboratories (Montgomery, TX) and used for immunofluorescence from BD Biosciences (Franklin Lakes, NJ). Horseradish peroxidase-conjugated anti-mouse and anti-rabbit immunoglobulin G (IgG) antibodies were from Chemicon (Billerica, MA). Cy3-conjugated anti-mouse IgG antibody and Alexa-Fluor 488 anti-rabbit antibody were from Jackson ImmunoResearch Laboratories (West-Grove, PA). Anti- β -actin monoclonal antibody was purchased from Sigma-Aldrich. Anti-HA monoclonal antibody was purified from the 12CA5 hybridoma. The anti-p85 antibody has been described (28).

Plasmids

The Afadin cDNA constructs pEGFP-N2-AF6i3-Myc and pEGFP-N2-AF6i1-Myc were a gift from Mihaela Lorguer and have been described (24). ShRNA rescue mutants were generated by introduction of 6 silent mutations using the following primer: 5' GGA ACG CCA GCG TCT TTT TTC ACA AGG ACA GGA CGT CTC TAA TAA AGT GAA AGC TTC TCG 3'. The shRNA resistant mutants with phosphorylation site mutations were generated by site

directed mutagenesis using the following primers: S1718A: 5' GAA CGC CAG CGT CTT
 TTT GCA CAA GGA CAG GAC G 3'; S1718D: 5' ACA TTC AAG GAA CGC CAG CGT CTT
 TTT GAT CAA GGA CAG GAC GTC 3'; S1718E: 5' ACA TTC AAG GAA CGC CAG CGT
 CTT TTT GAG CAA GGA CAG GAC GTC 3'. All sequences were verified by sequencing.
 pcDNA3-Myr-HA-Akt1, pcDNA3-Myr-HA-Akt2, pcDNA3-Myr-HA-Akt3 from William Sellers
 (Addgene plasmids 9008, 9016, 9017) (29). HA-GSK3 β has been described (30). JP1520-
 GFP, JP1520-PIK3CA-WT-HA, JP1520-PIK3CA-H1047R-HA, JP1520-PIK3CA-E545K-HA
 were from Joan Brugge (Addgene plasmids 14570, 14571, 15572) (31).

RNA Interference

For shRNA-silencing, a set of single-stranded oligonucleotides encoding the Akt1 or Akt2
 target shRNA and its complement were previously described(11). Akt3, sense, 5' CCG GCT
 GCC TTG GAC TAT CTA CAT TCT CGA GAA TGT AGA TAG TCC AAG GCA GTT TTT G
 3'; Akt3 antisense, 5' AAT TCA AAA ACT GCC TTG GAC TAT CTA CAT TCT CGA GAA
 TGT AGA TAG TCC AAG GCA G 3' (Sigma-Aldrich). For silencing of Afadin, specific
 sequences for l-Afadin were used: shAfadin #2, sense, 5' CCG GAA GGT CAA GAT GTA
 TCC AAT ACT CGA GTA TTG GAT ACA TCT TGA CCT TTT TTT G 3'; shAfadin #2,
 antisense: 5' AAT TCA AAA AAA GGT CAA GAT GTA TCC AAT ACT CGA GTA TTG GAT
 ACA TCT TGA CCT T 3'; shAfadin #3, sense: 5' CCG GAA ACT TGA CAT TCA AGG AAC
 GCT CGA GCG TTC CTT GAA TGT CAA GTT TTT TTT G 3'; shAfadin #3, antisense: 5'
 AAT TCA AAA AAA ACT TGA CAT TCA AGG AAC GCT CGA GCG TTC CTT GAA TGT
 CAA GTT T 3'. The oligonucleotide pair for each target was annealed and inserted into
 pLKO. To produce lentiviral supernatants, 293T cells were co-transfected with control or
 shRNA-containing pLKO vectors, VSVG, and psPAX2 for 48 hr.

182 **In Vitro Kinase Assays**

183 HeLa cells were transfected with Myc-Afadin-wild type or Myc-Afadin-S1718A. 24 hr after
 184 transfection, cells were serum starved for 16 hr. Afadin was immunoprecipitated from cell
 185 extracts and incubated with 500 ng recombinant Akt1 or Akt2 (Cell Signaling Technology) in
 186 the presence of 250 μ M cold ATP in a kinase buffer for 1 hr at 30°C. The kinase reaction
 187 was terminated by addition of SDS-PAGE sample buffer.

188

189 **Transwell migration assay**

190 Cells (1×10^5) in serum-free medium containing 0.1% BSA were added to upper chambers
 191 of transwell filters (8 μ m pore size; BD Biosciences) in triplicate. NIH 3T3 -cell-conditioned
 192 medium, or growth medium from MCF10A was added to lower chambers. After 2–16 hr
 193 incubation at 37°C, non-migrated cells were removed and cells that had migrated to the
 194 bottom of the filters were fixed and stained using the Hema-3 stain set Protocol (Fisher
 195 Scientific; Pittsburgh, PA).

196

197 **Immunofluorescence**

198 Cells plated on coverslips were fixed with 2% paraformaldehyde for 10 min, permeabilized
 199 with 0.5% Triton X-100 and blocked with 1% BSA in 20 mM Tris-HCl (pH 7.5) for 20 min,
 200 then incubated with antibodies for 1 hr (anti-Afadin, anti-pAfadin, anti-pAkt S473, anti-E-
 201 cadherin, anti-CENP-A, anti-NuP98, anti-Fibrillarin, anti-Histone H3; 1:200). After washing
 202 twice with phosphate-buffered saline (PBS), cells were incubated with Cy3-conjugated anti-
 203 mouse IgG antibody or Alexa-fluor 488 anti-Rabbit IgG for 1 hr. F-actin was visualized with
 204 Alexa Fluor 647-conjugated phalloidin (Life Technologies; Grand Island, NY). Cells were
 205 then rinsed twice with PBS and mounted with Prolong Gold antifade reagent-4,6-diamidino-

2-phenylindole (DAPI) (Life Technologies). Images of cells were acquired using a fluorescence microscope (Nikon Eclipse Ti) and digital image analysis software (NIS-Elements, Nikon). Magnification plan Apo VC 60x/1.40 oil. Experiments to determine staining with phospho-Afadin S1718 antibody always included co-staining with total Afadin antibody in order to ensure specificity of staining.

Immunoblotting and Immunoprecipitation

Cells were lysed in RIPA as previously described (13). Lysates were resolved on 6-10% acrylamide gels by SDS-PAGE and transferred to PVDF membrane (EMD Millipore; Billerica MA). The blots were blocked in TBST buffer (10 mM Tris-HCl [pH 8], 150 mM NaCl, 0.2% Tween 20) containing 5% (w/v) non-fat dry milk for 30 min and then incubated with the specific primary antibody diluted in blocking buffer at 4°C for 16 hr. Membranes were washed three times in TBST and incubated with horseradish peroxidase-conjugated secondary antibody for 1 hr at room temperature. Membranes were washed three times and developed using enhanced chemiluminescence substrate (EMD Millipore). For immunoprecipitation, lysates were incubated with 1–2 µg antibody for 2–4 hr at 4°C followed by incubation with 15 µl protein A/G Sepharose beads (Amersham Biosciences; Pittsburgh). Immune complexes were washed with NETN buffer (0.5% NP-40, 1 mM EDTA, 20 mM Tris-HCl [pH 8], 100 mM NaCl). Precipitates were resolved by SDS-PAGE.

Subcellular Fractionation

Cells were fractionated using the Subcellular protein fractionation kit for cultured cells (Thermo Fisher Scientific; Rockford, IL) according the manufacturer's instructions.

229

230 **Tissue Microarrays**

231 Tissue microarrays (TMA) containing normal breast tissue (2 cores per case) and invasive
 232 breast cancer (2 cores per case) were constructed from archival FFPE breast tissue
 233 specimens obtained from Beth Israel Deaconess Medical Center under an institutionally-
 234 approved IRB protocol for discarded de-identified tissues. Double immunofluorescence for
 235 Afadin and E-cadherin was performed. Antigen retrieval was performed by boiling the slides
 236 for 10 min in 10mM sodium citrate pH6 with a pressure cooker. The sections were then
 237 incubated with 1mg/ml sodium borohydride (MP Biomedicals; Solon, OH) for 5 min at room
 238 temperature. Sections were incubated with 5% normal donkey serum (Jackson
 239 ImmunoResearch) for one hr at room temperature. Slides were then incubated with mouse
 240 anti-Afadin (1:100, BD Biosciences) and Rabbit anti-E-Cadherin (1:100, Cell Signaling
 241 Technology) overnight at 4°C. The slides were washed and incubated with Alexa 488
 242 conjugated Donkey anti-rabbit or anti-mouse secondary antibodies (Jackson
 243 ImmunoResearch Lab, 1:200). Samples were then washed and mounted with Prolong Gold
 244 anti-fade mounting media containing DAPI (Invitrogen). We digitally acquired 98
 245 microscopic images of normal breast tissue and 98 microscopic images of invasive breast
 246 cancer at 63 X magnification. Features were extracted from the digital images in ImageJ
 247 and statistical analyses were performed using Jython and R. To determine the statistical
 248 significance of the difference in Afadin nuclear localization score in normal breast as
 249 compared with invasive breast cancer we performed a two-sided Student's T-test.

250

251 Results

252 Akt phosphorylates Afadin at Ser1718

253 Global phosphoproteomic analyses have revealed that the adherens junction protein
 254 Afadin is phosphorylated at serine 1718 (Ser1718) (16), in a sequence within the actin-
 255 binding domain that conforms to the optimal Akt consensus motif RXRXXS/T (32). The motif
 256 surrounding Ser1718 is evolutionarily conserved from *Drosophila* to mammals (Fig. 1A).
 257 Since the PI 3-K/Akt pathway modulates all phenotypes associated with breast cancer and
 258 does so by phosphorylating substrate proteins to transduce the signal, we evaluated Afadin
 259 protein expression in breast cancer cell lines. Afadin is highly expressed in various breast
 260 cancer cell lines, including basal and luminal molecular subtypes as well as the non-
 261 tumorigenic line MCF10A (Fig. 1B).

262 To determine whether Afadin is a substrate of Akt, MCF10A (Fig. 1C) and HeLa cells
 263 (Supplementary Fig. S1A) were serum-starved and stimulated with IGF-1. Stimulation leads
 264 to phosphorylation of Afadin at Ser1718 as detected by a phospho-specific anti-pSer1718
 265 antibody. Ser1718 phosphorylation induced by IGF-1 is significantly inhibited by wortmannin
 266 (a pan-PI 3-K inhibitor), BEZ-235 (a dual PI 3-K and TORC1 inhibitor), A66 (a p110 α
 267 specific inhibitor) and MK2206 (an allosteric pan-Akt inhibitor) (33-36) (Fig. 1C). By contrast,
 268 Ser1718 phosphorylation is not blocked by Rapamycin (an mTOR inhibitor), GSK650394
 269 (an SGK (serum and glucocorticoid-induced kinase) inhibitor), PF4708671 (an S6K1 (p70
 270 S6-kinase-1) inhibitor) or CGK733 (an ATM/ATR (Ataxia Telangiectasia Mutated/ ATM-
 271 related (ATR)) inhibitor) (37-41). Akt phosphorylates Afadin specifically at Ser1718 since a
 272 Ser1718Ala (S1718A) mutant is not phosphorylated in response to IGF-1, and no additional
 273 Akt consensus motifs are found in the Afadin amino acid sequence. Moreover, the short
 274 isoform of Afadin (s-Afadin) is not phosphorylated in response to IGF-1, (Fig. 1D), consistent
 275 with the fact that it lacks the Ser1718 motif (Fig. 1A)

To determine whether one or more Akt isoform can phosphorylate Afadin in cells, Akt1, Akt2 and Akt3 were silenced using specific shRNA's introduced into MCF10A cells (Fig. 2A) and HeLa cells (Supplementary Fig. S1B). Silencing individual Akt isoforms partially attenuates Ser1718 phosphorylation, whereas combination silencing of Akt1, Akt2 and Akt3 leads to a complete abrogation of the pSer1718 signal (16% in pLKO versus 100% in pLKO cell stimulated with IGF-1; Akt1 silencing (104%), silencing Akt2 (45%), silencing Akt3 (25%) and combined Akt1/Akt2/Akt3 silencing (34%), normalized relative to total Afadin). Moreover, co-expression of constitutively active, myristoylated Akt1, Akt2 and Akt3 alleles leads to enhanced Afadin Ser1718 phosphorylation (Fig. 2B), and purified recombinant Akt1 or Akt2 can directly phosphorylate Afadin at Ser1718 in an *in vitro* protein kinase assay (Fig. 2C). Similarly, co-expression of the oncogenic *PIK3CA* alleles H1047R and E545K that stimulate hyperactivation of Akt also induces Afadin Ser1718 phosphorylation in MCF10A cells (Fig. 2D) and HeLa cells (Supplementary Fig. S1C). In aggregate, these data demonstrate that Afadin is phosphorylated by all Akt isoforms downstream of PI 3-K, but is not a substrate for other AGC kinases including SGKs and S6K1.

Phosphorylation of Afadin at Ser1718 promotes nuclear localization

Since Afadin is an adherens junction protein, we next evaluated the consequence of Ser1718 phosphorylation by Akt on cellular localization. Using immunofluorescence of IGF-1-stimulated MCF10A cells, we detect phosphorylated, activated Akt (pS473) within 20 min of stimulation (Supplementary Fig. S2). Under these conditions, total Afadin shows a predominantly membrane-restricted localization. However, within 1 hr of stimulation, Afadin membrane localization is significantly diminished, concomitant with the appearance of nuclear localization as evidenced by co-staining with DAPI (Fig. 3A and Supplementary Fig. S2). Nuclear localization of Afadin is most evident by 6 hr post-stimulation, with a punctate

nuclear staining pattern (Fig. 3A, IGF-1, 6hr). Nuclear translocation of total Afadin is dependent on PI 3-K and Akt activity, since wortmannin, MK2206, A66 and BEZ235 block nuclear localization in favor of membrane localization (Fig. 3A and Supplementary Fig. S3A). Quantification of the nuclear translocation in response to IGF-1 and Akt inhibitor is depicted in the bar graph (Fig. 3A). The pSer1718 antibody also reveals Afadin nuclear localization in response to IGF-1 stimulation (Fig. 3B, quantification shown in the corresponding bar graph).

To explore the contribution of Ser1718 in plasma membrane to nucleus translocation, an shRNA silencing and rescue experiment was performed. MCF10A cells were transduced with Afadin shRNA and non-silenceable wild-type, Ser1718Ala (S1718A) and Ser1718Glu (S1718E) mutants transiently expressed. As predicted, in full serum conditions, wild-type Afadin is localized to the plasma membrane and nucleus, Ser1718Ala Afadin is localized predominantly to the plasma membrane, whereas the phosphomimetic Ser1718Glu mutant shows an exclusively nuclear localization (Fig. 4A, quantification shown in the corresponding bar graph). Moreover, co-expression of constitutively activated, myristoylated Akt1, Akt2 or Akt3 alleles also promotes Afadin nuclear localization compared to control cells (Fig. 4B and corresponding bar graph, and Supplementary Fig. S3B). Therefore, Akt signaling promotes the relocalization of Afadin from the plasma membrane to the nucleus in a manner that depends on Ser1718 phosphorylation.

To further explore the mechanism of Afadin nuclear translocation, cell fractionation was performed. In agreement with the immunofluorescence data, IGF-1 stimulation of MCF10A cells results in a time-dependent decrease of total Afadin from the cytoplasm and membrane compartments, concomitant with an increase of Afadin in the nuclear compartment, most dramatically evident 6 hr post-stimulation (Fig. 5A, left panels). Treatment with the Akt inhibitor MK2206 attenuates this translocation (Fig. 5A, right panels). We also evaluated the consequence of Afadin phosphorylation on protein stability. Serum-

starved cells were stimulated over time with IGF-1 in the presence or absence of the protein synthesis inhibitor cycloheximide as well as Akt inhibitor. As observed in Fig. 5B, prolonged treatment of cells with MK2006 results in a reduction of total Afadin expression, which is further enhanced in the presence of cyclohexamide. Similar results are observed when the same cells are examined by immunofluorescence (Fig. 5C and corresponding bar graph). These data indicate that Akt signaling, in addition to promoting nuclear relocalization, also promotes stabilization of Afadin.

Phosphorylation of Afadin at Ser1718 enhances migration and perturbs cell-to cell adhesion

We next reasoned that since Akt promotes relocalization of Afadin from adherens junctions to the nucleus, this would likely have a profound impact on cell adhesion and cell migration, phenotypes that are dependent on intact adherens junctions. In this context, previous studies have shown that Afadin shRNA enhances migration of MCF7, SK-BR3 and MDA-MB-231 cells (27). Yet in other studies Afadin silencing reportedly enhances cell adhesion in T cells (42). The contribution of Afadin to cell migration is therefore likely to be context dependent. In BT549 and MDA-MB-468 breast cancer cells, that express high levels of Afadin and exhibit PI 3-K pathway activation due to *PTEN* inactivation and consequently predominantly nuclear localized Afadin (Supplementary Fig S5A and S5B), silencing Afadin with specific shRNA leads to a profound inhibition of cell migration (Supplementary Fig. S5C). Conversely, expression of wild-type or phospho-mimetic Ser1718Asp (S1718D) or Ser1718Glu (S1718E) Afadin in T47D cells, that do not express Afadin, leads to enhanced cell migration in a manner that is not phenocopied by the Ser1718Ala mutant (Fig. 6A). The cellular localization of these mutants expressed in T47D cells is in agreement with the localization observed in MCF10A cells (Supplementary Fig. S6 compared to Fig. 4A).

Consistent with the finding that Afadin promotes cell migration of breast cancer cells, silencing Afadin in MCF10A cells profoundly blocks migration in a manner that is partially rescued by re-expression of wild-type and Ser1718Asp (S1718D), but not Ser1718Ala (S1718A) mutants (Fig. 6B). Taken together, these results demonstrate that Afadin promotes breast cancer cell migration in a manner that depends, at least in part, on Ser1718 phosphorylation mediated by Akt.

Finally, since Afadin is localized to adherens junctions (20), we evaluated the consequence of Afadin relocalization on cell to cell adhesion using E-cadherin staining measured by immunofluorescence. In control serum starved MCF10A cells, both Afadin and E-cadherin show restricted membrane localization with defined cell to cell adhesion (Fig. 6C). By contrast, in cells transduced with Afadin shRNA, E-cadherin staining is significantly disrupted. A similar phenotype is observed in cells in which Afadin is silenced and the nuclear localized Ser1718Asp (S1718D) mutant is re-expressed (Fig. 6C, controls of wild type Afadin, S1718A and S1718E Afadin are shown in Supplementary Fig. S7). We conclude that membrane-localized Afadin is required for maintaining intact adherens junctions and productive cell to cell adhesion, such that loss of membrane localization and nuclear relocalization disrupts adhesion, concomitant with an increase in cell migration.

In order to address the specific nuclear compartment that Afadin localizes to, we performed co-localization experiments using a number of established nuclear markers: CENP-A, a centromere marker; NuP98, a nuclear envelope marker; Fibrillarin, a nucleolar marker; and Histone H3, a nucleosome or chromatin marker (Fig. 6D). The punctate nuclear pattern of Afadin does not colocalize with any of these markers. Future studies will address the specific nuclear compartment that Afadin localizes to, and the importance of this localization for the nuclear function of Afadin.

376 **Afadin localization in breast cancer**

377 We next assessed Afadin localization in human breast cancer. Tissue microarrays
 378 containing normal breast epithelium and invasive breast cancer tissue obtained from
 379 archival pathology specimens from 49 patients were used for localization evaluated by
 380 immunofluorescence using Afadin and E-cadherin staining. The quantification protocol and
 381 analysis is summarized in Supplementary Fig. S8. We identified significantly increased
 382 nuclear localization of Afadin in invasive breast cancer as compared to normal breast, with
 383 53% higher nuclear localization in invasive breast cancer (mean Afadin nuclear localization
 384 score in normal = 0.019 vs. mean Afadin nuclear localization score in cancer = 0.029; p,
 385 0.02). Fig. 7 shows representative images of normal breast tissue and invasive breast
 386 cancer specimens. From these data we conclude that the nuclear localization of Afadin,
 387 regulated by phosphorylation at Ser1718 by the Akt pathway, is clinically relevant for breast
 388 cancer progression.
 389

390 Discussion

391 We have identified and characterized a new substrate of Akt, the adherens junction
 392 protein Afadin. This finding adds to the list of the over 200 currently identified substrates of
 393 Akt kinases that transduce the PI 3-K/Akt signal to a plethora of biological and
 394 pathophysiological responses, particularly in the context of cancer (43). We have shown that
 395 Akt phosphorylates Afadin at Ser1718 in a motif that is evolutionarily conserved, indicating
 396 that this phosphorylation has evolved to modulate a key biological event. We have shown
 397 that physiological signaling in non-tumorigenic MCF10A cells stimulated with IGF-1 leads to
 398 Afadin phosphorylation, and that in breast cancer cell lines harboring pathway mutations
 399 such as oncogenic *PIK3CA* and *PTEN* inactivation, Afadin is phosphorylated in an Akt-
 400 dependent manner. Moreover, although several other AGC kinases such as a S6K and
 401 SGKs have an overlapping optimal consensus phosphorylation motif to Akt, only Akt is
 402 capable of phosphorylating Afadin in cells. Although a number of Akt isoform-specific have
 403 recently been identified, Afadin does not appear to be an isoform-specific substrate, at least
 404 in the breast cancer cell lines tested here.

405 Since mutations in genes that encode proteins in the PI 3-K and Akt signaling pathway
 406 are among the most common and frequent in human cancers, especially breast cancer,
 407 there are currently numerous clinical trials targeting both PI 3-K and Akt for therapeutic
 408 benefit. Hyperactivation of Akt due to oncogenic *PIK3CA* mutations as well as amplification
 409 and somatic mutations in Akt genes are common events in breast cancer etiology, and have
 410 been shown to result in cell transformation and cancer progression using mouse models.
 411 While the mechanisms by which PI 3-K and Akt promote cell transformation are well
 412 understood, the mechanisms by which this pathway promotes cancer progression at the
 413 level of tumor dissemination, invasion and metastasis are not as well characterized. In this
 414 context, it is now well-established that Akt isoforms play differential roles in promoting breast

cancer cell invasion and metastasis, whereby Akt1 does not enhance metastasis or can actually function as an invasion and metastasis suppressor, yet Akt2 promotes invasive migration leading to metastatic dissemination (12,44). A role for Akt3 in breast cancer progression has yet to be defined. The mechanistic basis for the differential role of Akt isoforms in mediating breast cancer progression is likely complex, and involves both differential intracellular localization as well as accessibility to specific substrates that control cell migration and invasion. A number of such substrates have been identified, including the actin-bundling protein palladin, Girdin and ACAP1 (9,13,14). In this context, it is interesting to note that Ser1718 lies within a previously-identified actin-binding domain (18), and as such it is possible that Ser1718 phosphorylation may modulate actin binding to promote cell migration, though this remains to be determined. Regardless, identifying the specific mechanisms by which the Akt pathway controls the phosphorylation of substrates that mediate cell migration is critical for a complete understanding of the contribution of this pathway in cancer progression, and in turn the development of drugs to target Akt kinases therapeutically.

We have shown that phosphorylation of Afadin promote relocalization from adherens junctions to the nucleus. This is most evident when evaluating Afadin localization by immunofluorescence, whereby IGF-1 stimulation leads to a relocalization of total and pSer1718 Afadin from the plasma membrane to the nucleus (Fig. 3). Similarly, a Ser1718Ala non-phosphorylatable mutant is membrane restricted and moreover a phosphomimetic Ser1718Asp mutant is constitutively localized to punctate nuclear structures in breast epithelial cells (Fig. 4). Most importantly, this localization phenocopies cell migration, whereby the Ser1718Ala cannot rescue the deficit in Transwell migration induced by Afadin shRNA, whereas the Ser1718Asp mutant can (Fig. 6). We conclude that Afadin phosphorylation at Ser1718 by Akt promotes cell migration, concomitant with a

relocalization from the membrane to the nucleus. Although localization of Afadin to the nucleus is dependent on productive Akt signaling and was observed in all experiments and in many cells visualized by immunofluorescence, it was not a quantitative event observed in 100% of the cell population. This is not surprising, however, since previous studies have reported cell-to-cell variability with respect to the activation status of Akt in a population of cells as result of *PIK3CA* heterogeneity (45). The proposed model is a bimodal distribution of Akt activation that is an invariable characteristic of exponentially growing cells. Limiting Akt activity to only 20%–30% of cells in a population serves, according to this study, two related purposes: it prevents senescence and maintains suboncogenic levels of PI 3-K activity in large populations.

What is more surprising is the relocalization of an adherens junction protein from the membrane to the nucleus in response to a single posttranslational modification, most obviously identified by the localization pattern of the Ser1718Asp and Ser1718Glu mutant Afadin. However there is some precedent to Afadin nuclear localization, since the short form of Afadin, s-Afadin, has been shown to be a dual-residency protein that localizes to the nucleus and to the plasma membrane in a manner dependent on growth factor signaling (46). In this study l-Afadin was not detected in the nucleus, although this is likely due to distinct experimental conditions and antibodies used to detect Afadin localization. What specifically mediates the translocation of Afadin from adherens junctions to the nucleus remains to be defined, and likely involves a multi-step process of nuclear import and retention.

However nuclear translocation is associated with an increase in breast epithelial and cancer cell migration. There are likely to be several mechanisms by which changes in Afadin localization mediate cell migration, since knocking out Afadin using specific shRNA decreases cell migration. At the same time, expression of wild-type or Ser1718Asp mutant

Afadin that is nuclear-restricted robustly enhances cell migration (Fig. 6B), indicating the simple removal of Afadin from adherens junction is not the only mechanism that affords cell migration. How a nuclear localized Afadin promotes cell migration remains to be determined, but could involve the induction of a transcriptional program. Although there is no evidence that Afadin can function as a transcriptional co-activator, this is reminiscent of β -catenin, also a component of adherens junctions that upon Wnt signaling translocates to the nucleus and functions as a transcriptional co-activator for the TCF/LEF transcription factor complex, and that in turn initiates a range responses including the epithelial to mesenchymal transition (EMT) (47). Whether Afadin functions in a similar manner remains to be determined. However, it is intriguing to note that either knocking out Afadin with shRNA or expression of a nuclear restricted Afadin mutant (Ser1718Asp) results in significant disruption of E-cadherin staining at the membrane (Fig. 6C).

Interestingly, loss of Afadin has been suggested to be a marker of poor prognosis in breast cancer, such that loss of Afadin actually promotes cell migration of MCF7, MDA-MB-231 and SKBR3 cells as measured in non-directional wound healing assays (27). In our studies performed in Transwell assays, in MCF10A, BT549 and MDA-MB-468 specific Afadin shRNA suppresses cell migration towards chemoattractants, and this can be effectively rescued by introduction of wild-type or phosphomimetic Afadin alleles (Fig. 6B). Therefore, the specific contribution of Akt signaling to Afadin and in turn cell migration is likely to be highly context-dependent, including the level of Afadin expression as well as the genetic background, in particular PI 3-K/Akt pathway mutations.

In summary, our study identifies Afadin as a new substrate of Akt that mediates cell migration in a manner that is dependent on cellular localization. Moreover, we show that nuclear-localized Afadin is a feature of human tumors as evident from localization studies from tissue microarrays. We propose that the phosphorylation of Afadin, an adherens

490 junction protein that is traditionally thought to reside exclusively at cell to cell adhesions and
491 whose phosphorylation modulates cell migration, is a previously uncharacterized
492 mechanism by which the Akt pathway promotes cancer progression. In this context,
493 although Afadin was originally defined as a “tumor-suppressor-like” protein, it may also
494 serve to function as a “tumor-promoting” protein at least in situations of pathophysiological
495 PI 3-K and Akt signaling.

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499

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 504 **immunofluorescence):** S. Elloul

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509 **Study supervision:** A. Toker

510

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- 642

643 **Figure legends**

644

645 **Figure 1: Akt phosphorylates Afadin at Ser1718**

646 A. Schematic of s-Afadin and l-Afadin showing domains and putative Akt consensus
 647 phosphorylation site at Ser1718 in l-Afadin. The phosphorylation site is evolutionarily
 648 conserved. Numbers on the left indicate the amino acid residue of the phosphorylation site.
 649 RA1,2, Ras associating domains 1 and 2; FHA, Forkhead associating domain; DIL, dilute
 650 domain; PDZ, PDZ domain; Pro, proline rich domain. Figure adapted from ref. (24). B.
 651 Expression of Afadin in breast cancer cell lines and HeLa cells evaluated by immunoblotting
 652 with total Afadin antibody. β -actin served as loading control. C. MCF10A cells were serum
 653 starved, pre-treated with the indicated inhibitors and stimulated with IGF-1 for 20 min. Whole
 654 cell lysates were immunoblotted with the indicated antibodies. p85 served as loading
 655 control. D. HeLa cells were transfected with vector control (pEGFP-N1), wild-type Afadin,
 656 Myc-Afadin-Ser1718Ala (S1718A) or Myc-s-Afadin (Afadin short form), serum starved and
 657 stimulated with IGF-1 for 20 min. Myc immunoprecipitates were immunoblotted with pAfadin
 658 antibody, and whole cell lysates immunoblotted with the indicated antibodies. p85 served as
 659 loading control. Results are representative of at least three independent experiments.

660

661 **Figure 2: PI 3-K and Akt Signaling Promotes Afadin Phosphorylation in breast cancer** 662 **cells**

663 A. Akt1, Akt2 and Akt3 shRNA transduced alone or in combination in MCF10A cells. Cells
 664 were stimulated with IGF-1 and whole cell lysates immunoblotted with the indicated
 665 antibodies. B. MCF10A cells transfected with control vector or myristoylated Akt alleles
 666 (MyrAkt1,2,3), cells were serum starved for 16 hr and whole cell lysates immunoblotted with
 667 the indicated antibodies. C. In vitro kinase assay using wild type Afadin, Afadin Ser1718Ala

(S1718A) or control immunoprecipitated with Myc antibody. Akt substrate GSK-3 β expressed in HeLa cells and immunoprecipitated with HA antibody and served as a positive control. Precipitates were incubated with purified recombinant Akt1 or Akt2. Whole cell lysates were immunoblotted with the indicated antibodies. D. MCF10A cells infected with vector control, PIK3CA wild type, PIK3CA H1047R, PIK3CA E545K, cells serum starved and whole cell lysates immunoblotted with the indicated antibodies. In all cases p85 served as loading control. Results are representative of at least three independent experiments.

Figure 3: IGF-1 Stimulation Promotes Afadin Nuclear Localization

A. MCF10A cells were serum starved and stimulated with IGF-1 for 1 or 6 hr, either alone or following treatment with MK2206 (1 μ M). Immunofluorescence was performed with the indicated antibodies. Quantification of the nuclear staining is presented in the bar graph. (Student T-test, * $p < 0.05$, ** $p < 0.01$).

B. MCF10A cells serum starved or stimulated for 6 hr with IGF-1, and immunofluorescence performed with the indicated antibodies. In all cases nuclei were stained with DAPI. Images are representative of multiple fields, and of at least three independent experiments. Quantification of the nuclear staining is presented in the bar graph. (Student T-test, ** $p < 0.01$). Magnified single channel staining for Afadin is presented in Supplementary Fig S4.

Figure 4: Phosphorylation of Afadin Promotes Nuclear Relocalization

A. MCF10A cells infected with control, pLKO or Afadin shRNA, and transfected with wild-type (WT) Afadin, Ser1718Ala (S1718A) or Ser1718Glu (S1718E) and immunofluorescence performed with the indicated antibodies. Quantification of the nuclear staining of Afadin is presented in the graph (** $p < 0.01$, Student T-test). B. MCF10A cells transfected with

pcDNA3 or myristoylated Akt1 alleles and immunofluorescence performed with the indicated antibodies. In all cases nuclei were stained with DAPI. Quantification of the nuclear staining of Afadin is presented in the bar graph (* $P < 0.05$, Student T-test). Images are representative of multiple fields, and of at least three independent experiments. Magnified single channel staining for Afadin is presented in Supplementary Fig S4.

Figure 5: Afadin Phosphorylation Modulates Protein Stability

A. MCF10A cells serum starved for 18 hr and stimulated with IGF-1 for 3 or 6 hr. Alternatively, cells in full serum were treated with MK2206 for 3 or 6 hr. Cytoplasmic and nuclear extracts were prepared and immunoblotted with the indicated antibodies. B and C, MCF10A cells serum starved for 18 hr and stimulated for 3 or 6 hr with IGF-1 either alone or with 20 μ g/ml cycloheximide for 3 or 6 hr. Alternatively, cells in full serum were treated with MK2206 as indicated. Whole cell lysates were blotted with the indicated antibodies (B) or immunofluorescence was performed (C) Quantification of the nuclear staining of Afadin is presented in the bar graph (* $p < 0.05$, Student T-test). Results are representative of multiple fields and at least three independent experiments. Magnified single channel staining for Afadin is presented in Supplementary Fig S4.

Figure 6: Afadin Phosphorylation Promotes Cell Migration

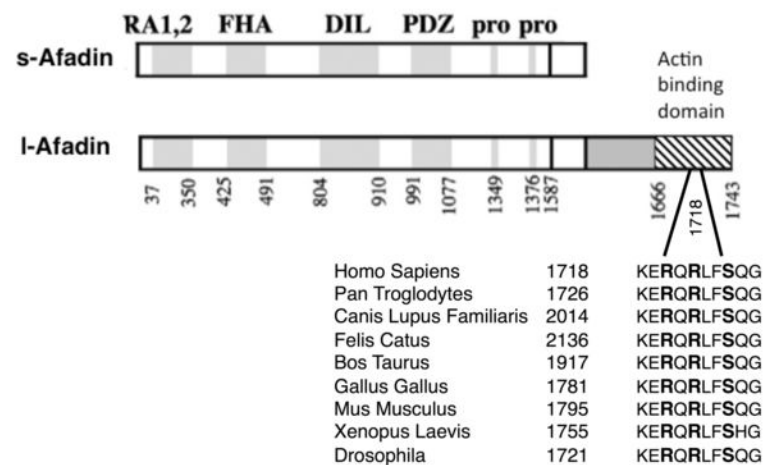
A. T47D transfected with wild-type (WT) Afadin, Ser1718Ala (S1718A), Ser1718Asp (S1718D) or Ser1718Glu (S1718E) Afadin. Transwell cell migration was assessed. In all cases data are represented as mean \pm SEM. * $p < 0.05$, ** $p < 0.01$ (Student T-test). B. MCF10A cells infected with Afadin shRNA and transfected with wild-type (WT), Ser1718Ala (S1718A) or Ser1718Asp (S1718D) mutants. Transwell cell migration was evaluated. In all cases data are represented as mean \pm SEM. ** $p < 0.01$ (Student T-test). C. MCF10A cells

transduced with shRNA with or without transfection of Afadin Ser1718Asp (SD) were evaluated by immunofluorescence with the indicated antibodies. Results are representative of multiple fields and at least three independent experiments. D. Immunofluorescence of MCF10A cells was performed on growing cells using the indicated nuclear markers antibodies (green channel), Afadin antibodies (red channel) and DAPI staining (blue channel). Results are representative of multiple fields and three independent experiments.

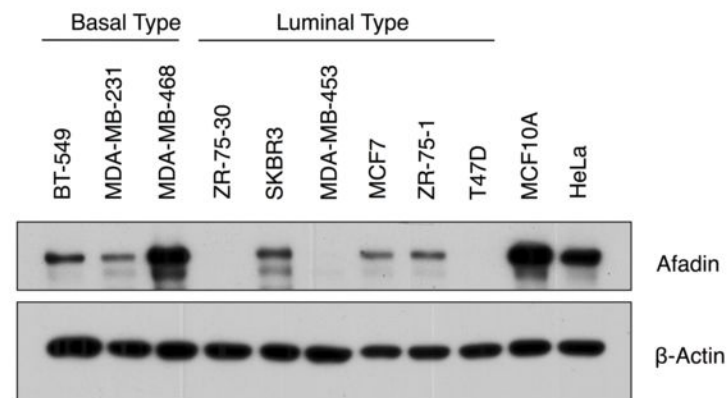
Figure 7: Afadin Localization in Breast Cancer

Tissue microarrays containing normal and invasive breast cancer from surgical specimens were subjected to immunofluorescence with the following antibodies: Afadin (red channel), E-cadherin (green channel) and DAPI (blue channel). Digital images were acquired with 63 X magnification. Representative images from normal tissue and invasive breast cancer are shown.

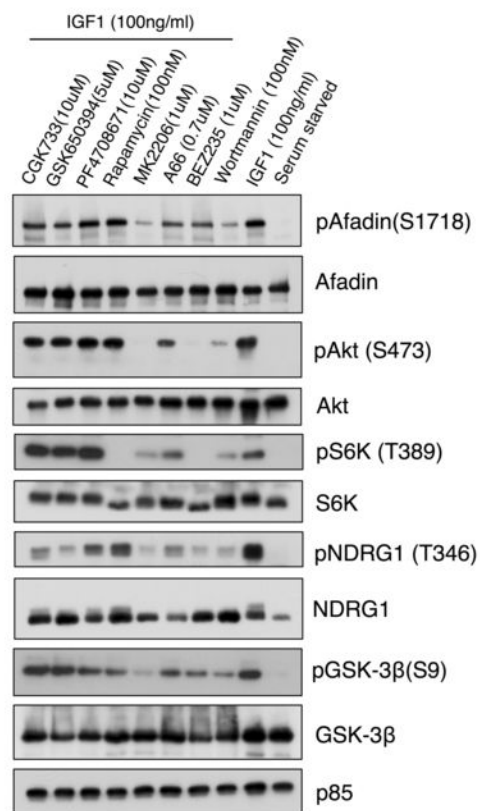
A



B



C



D

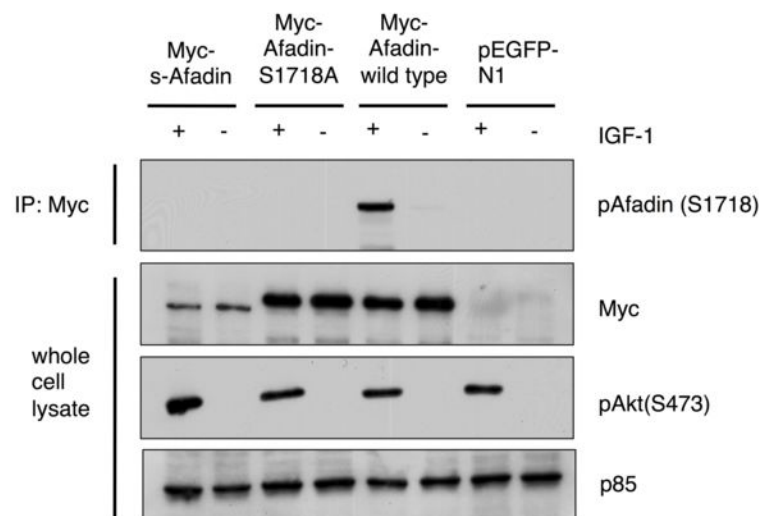


Fig. 1

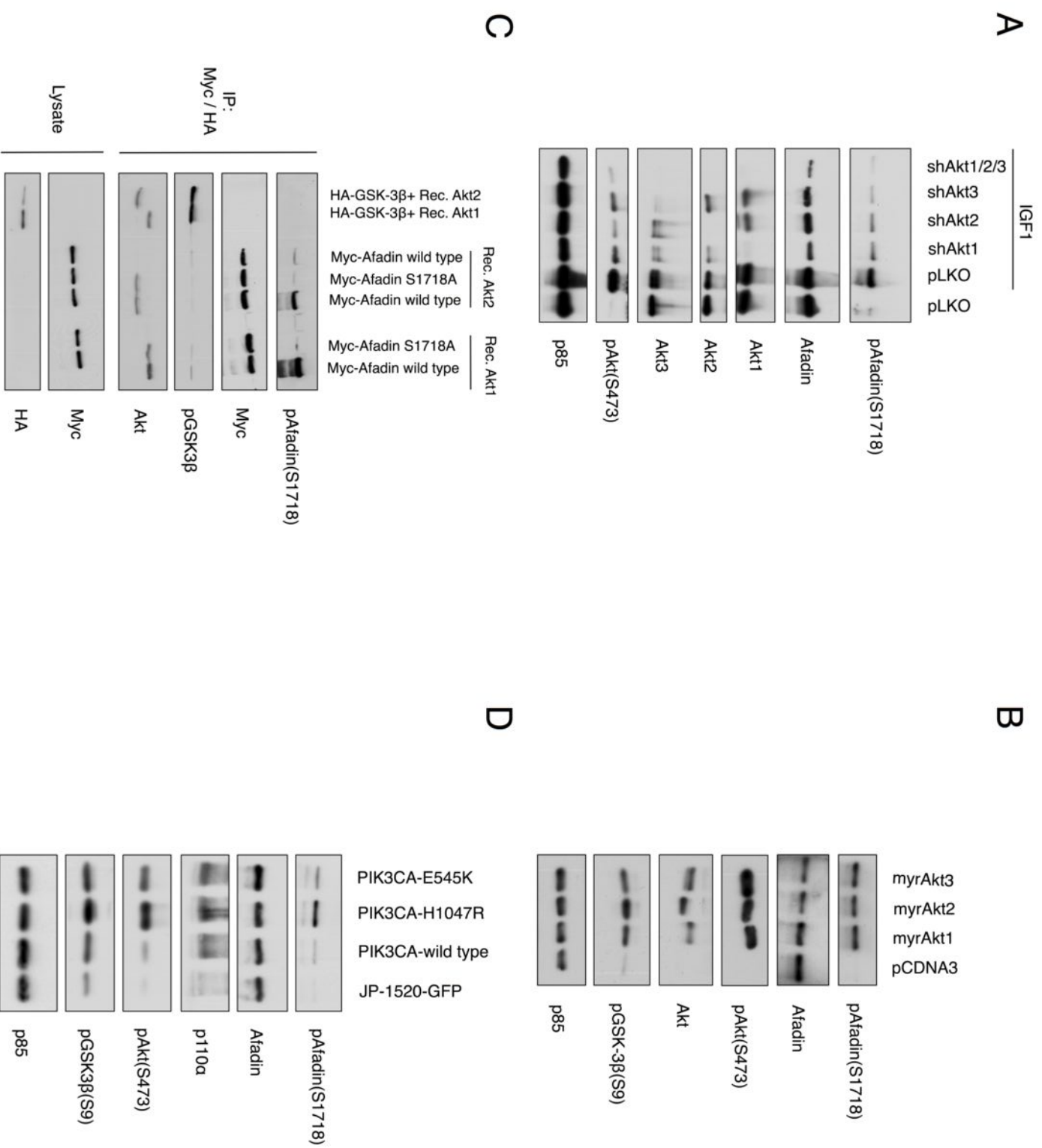


Fig. 2

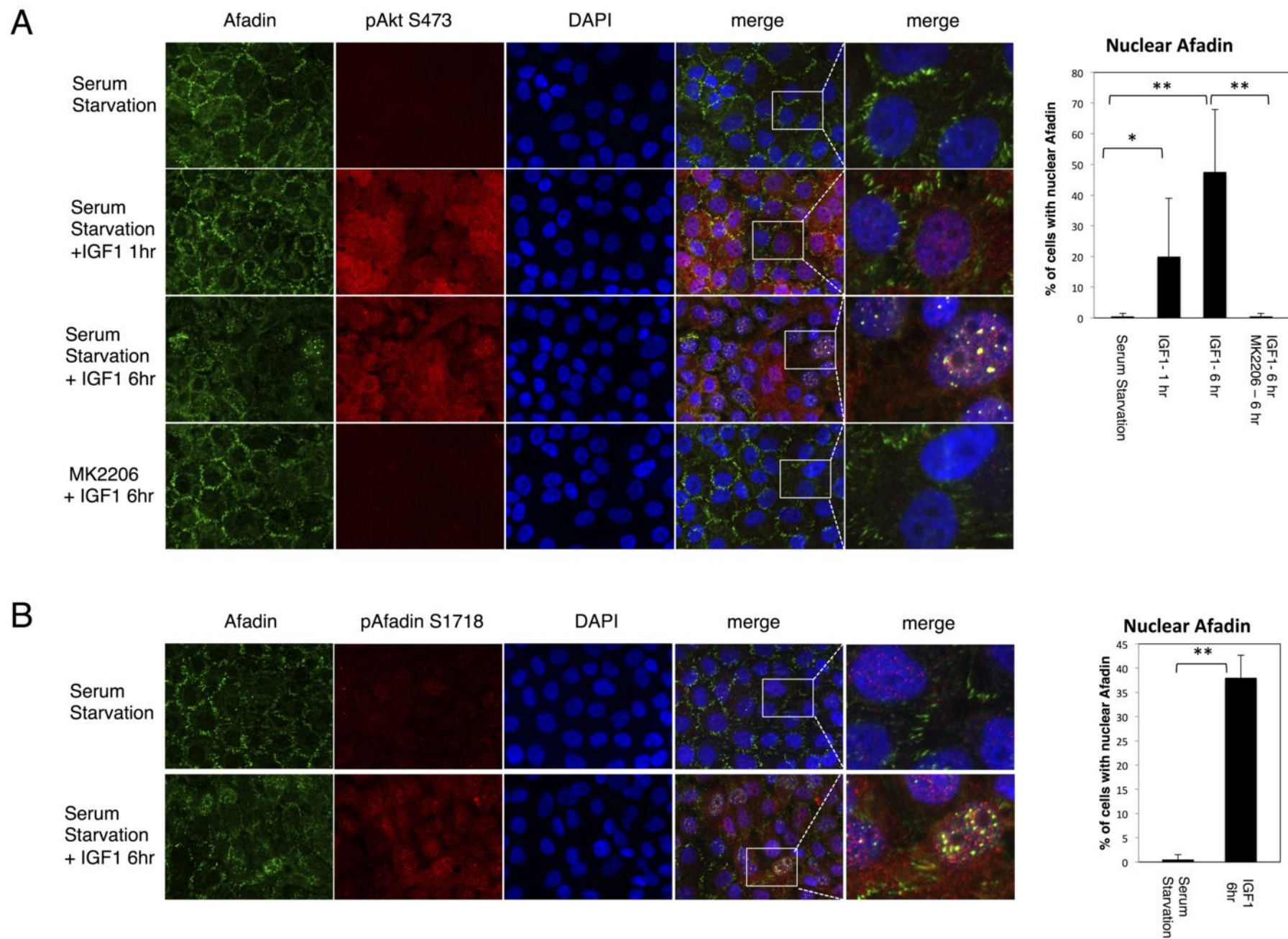


Fig. 3

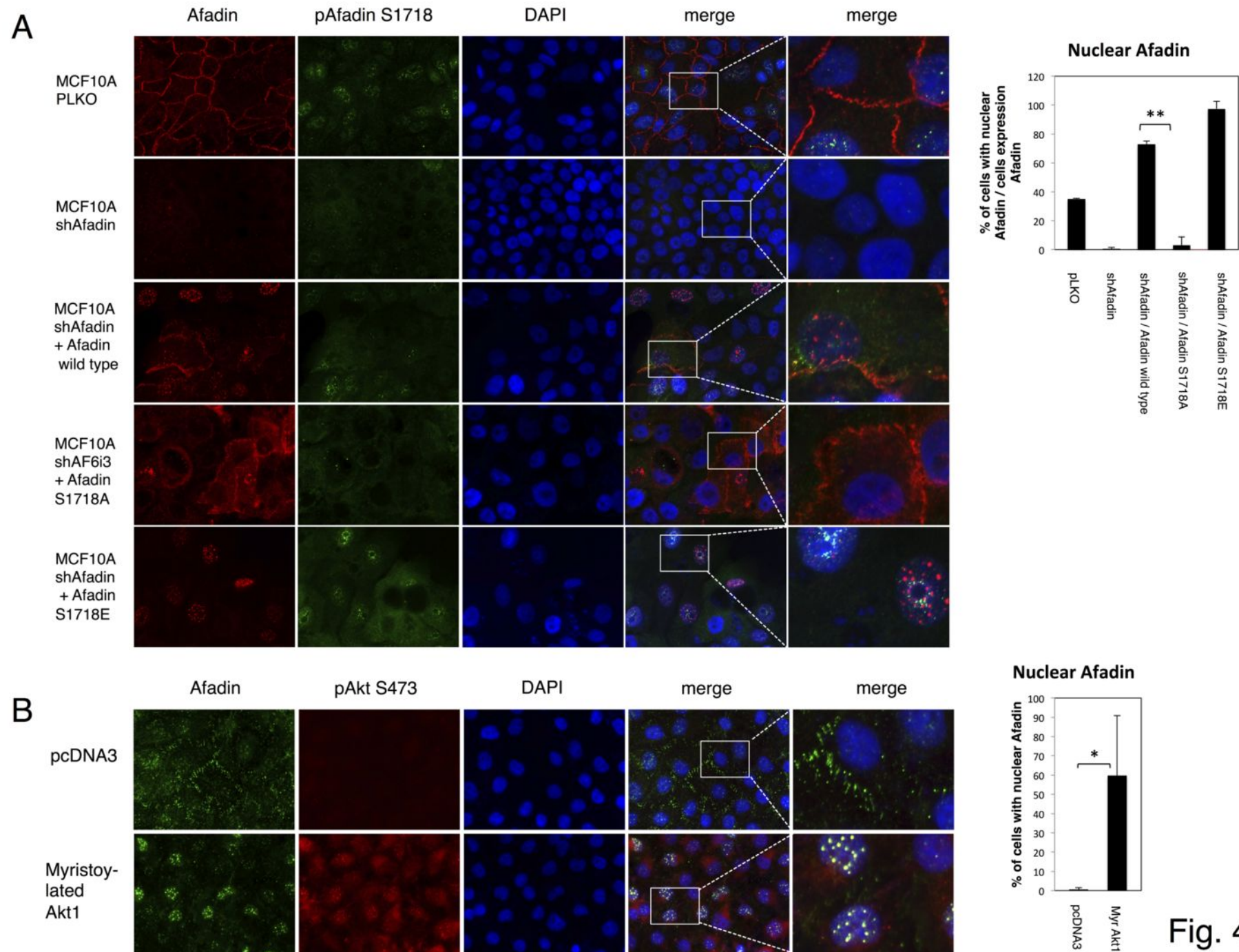
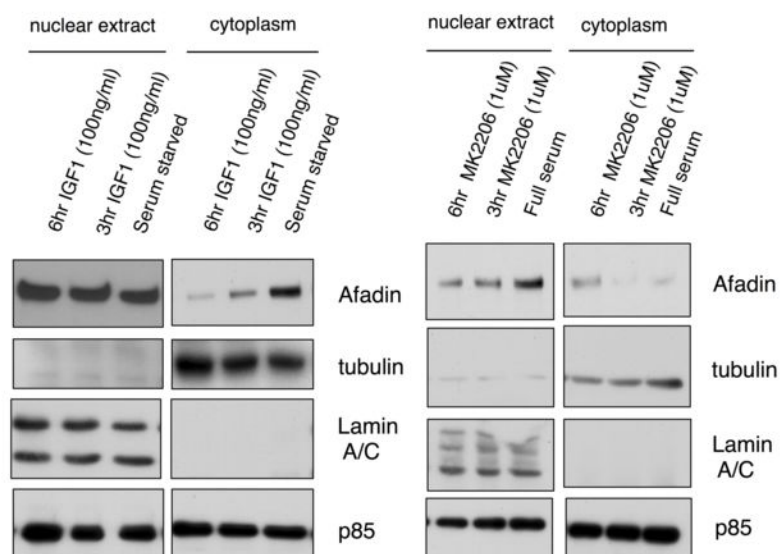
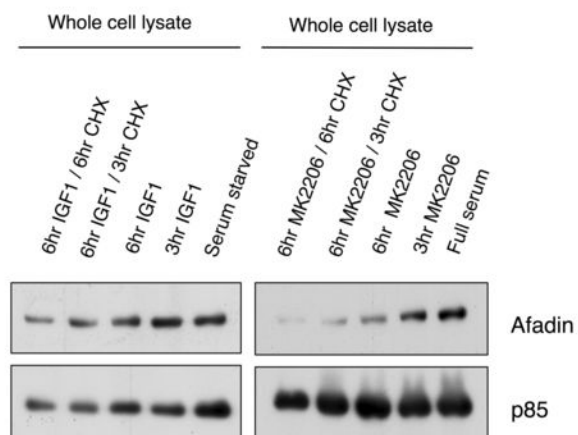


Fig. 4

A



B



C

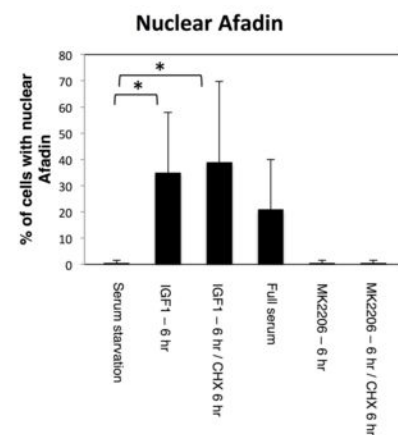
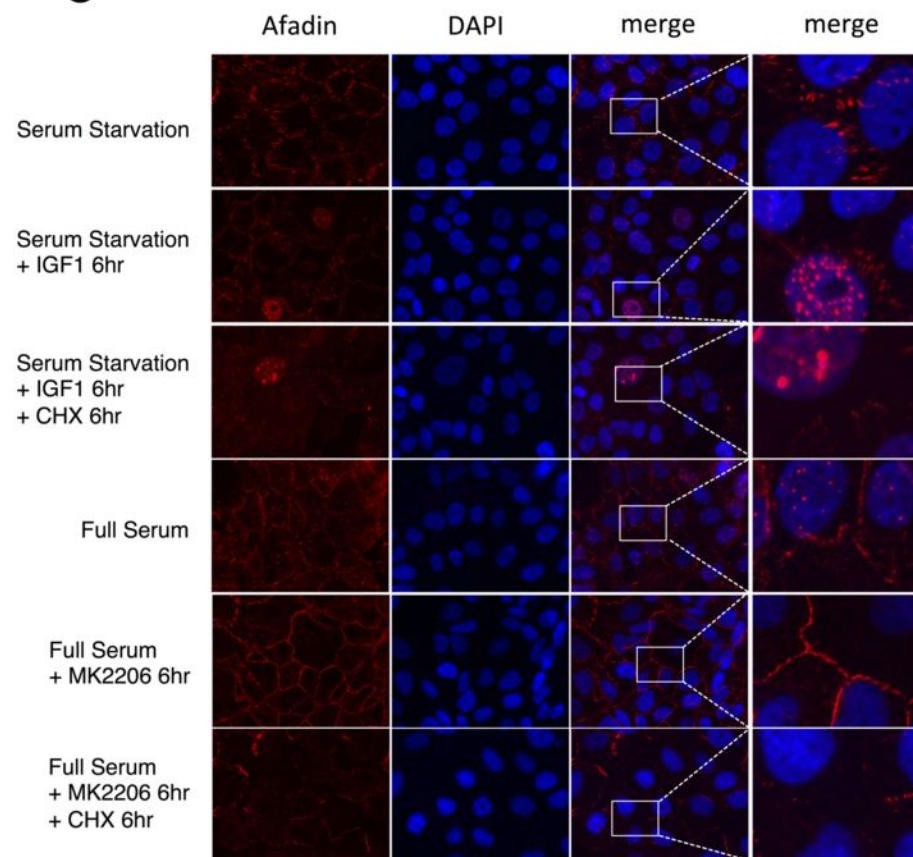


Fig. 5

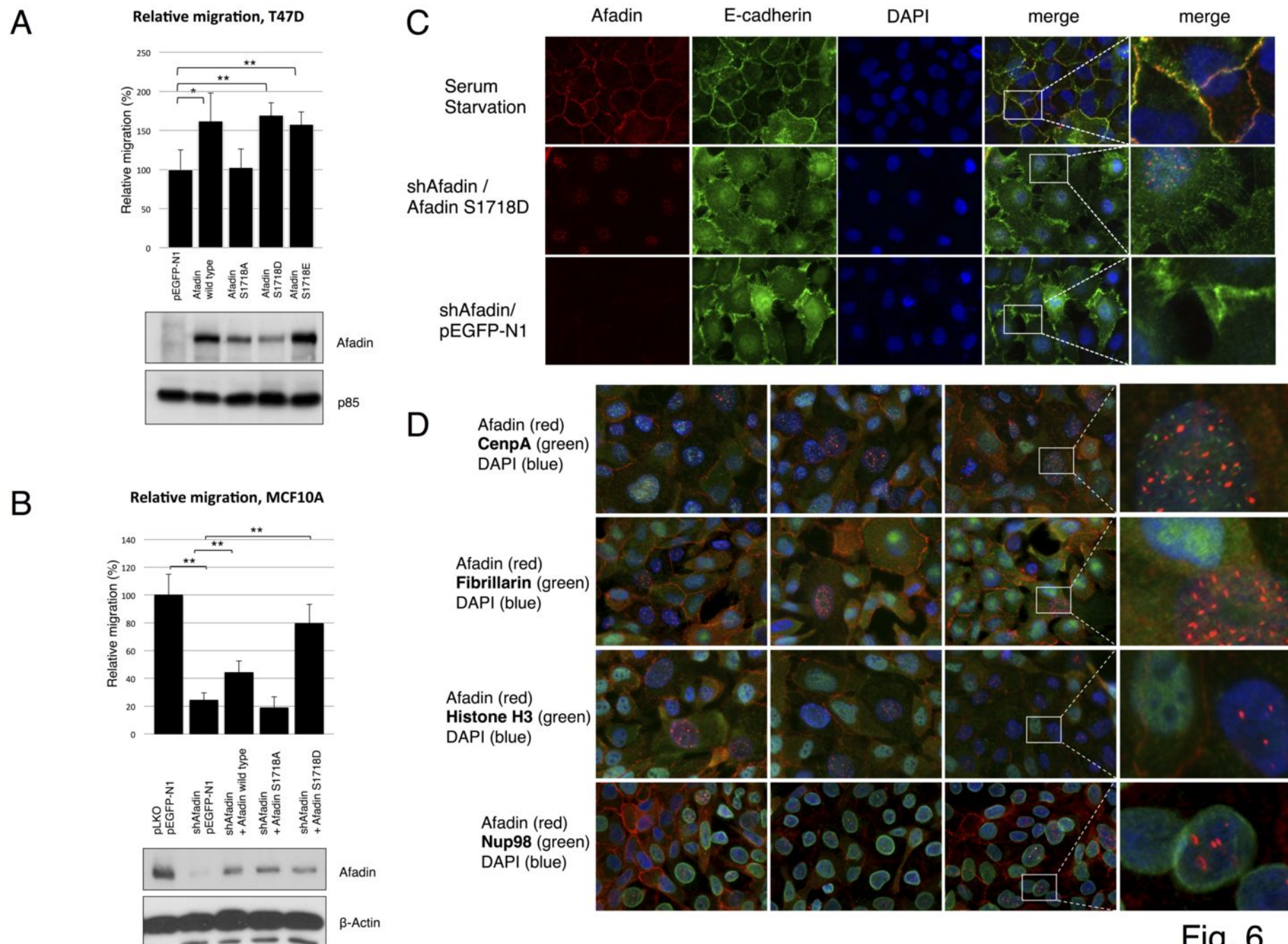
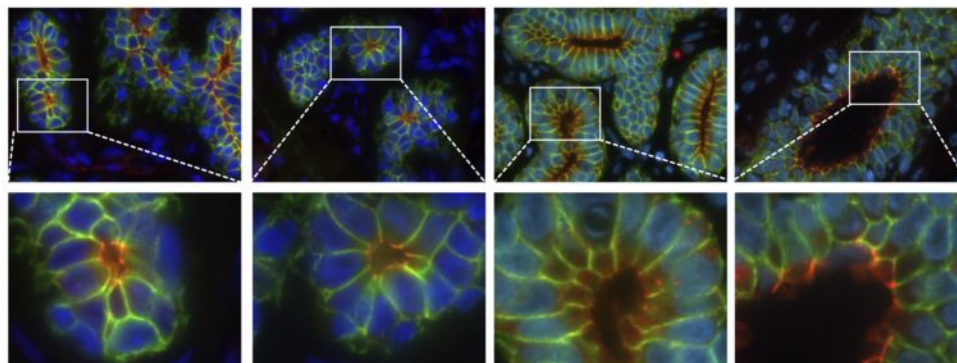


Fig. 6

Normal breast
tissue



Invasive breast
carcinoma

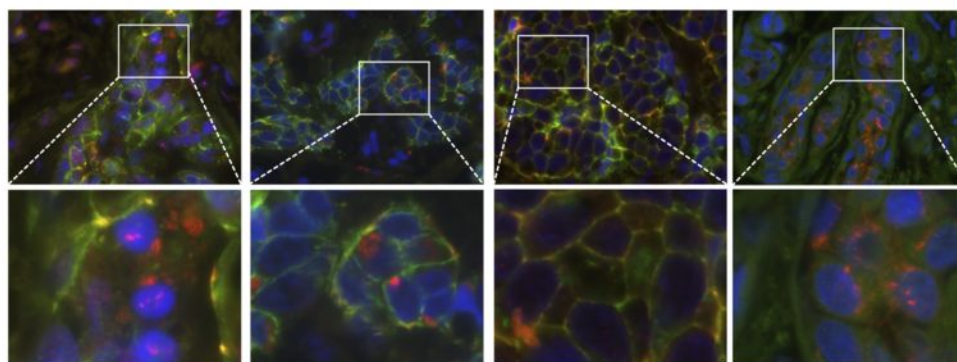
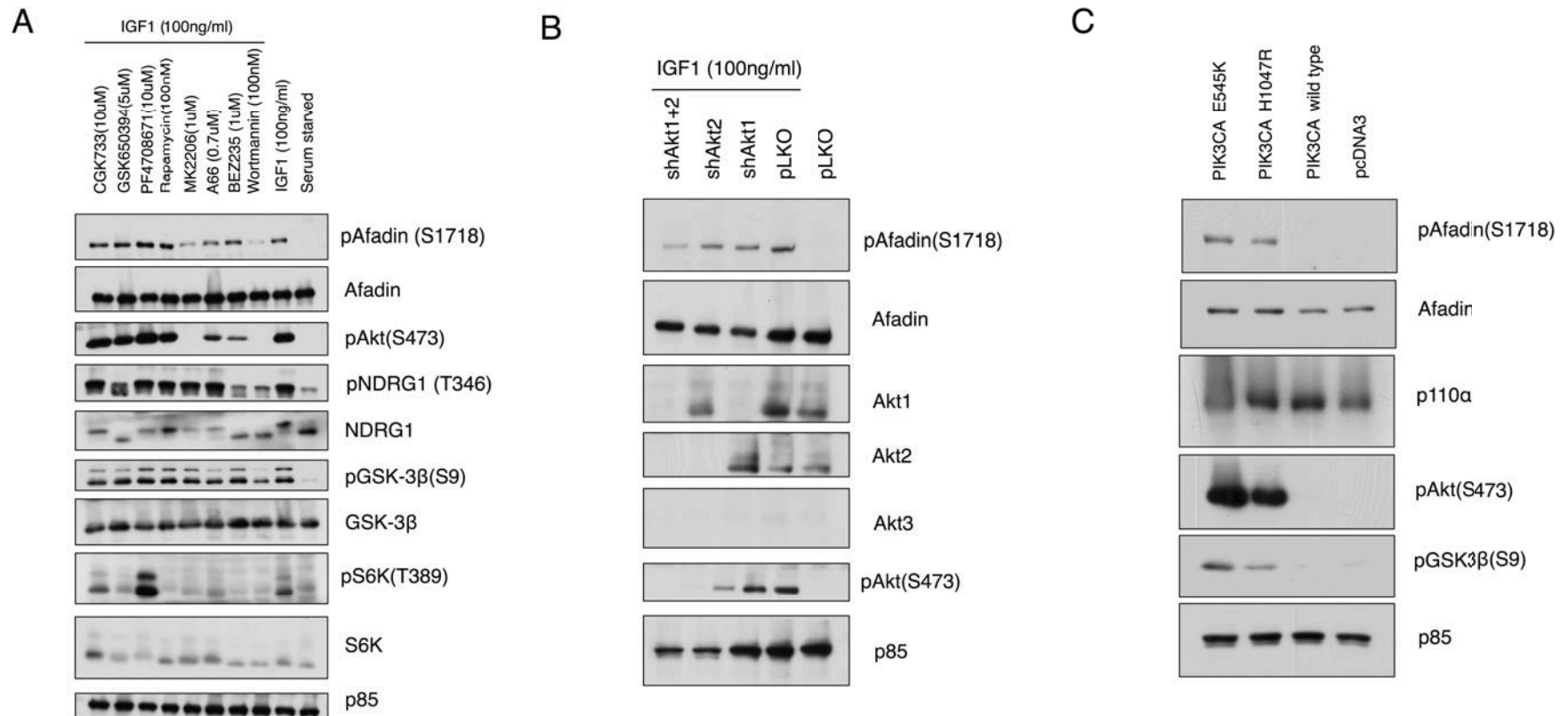


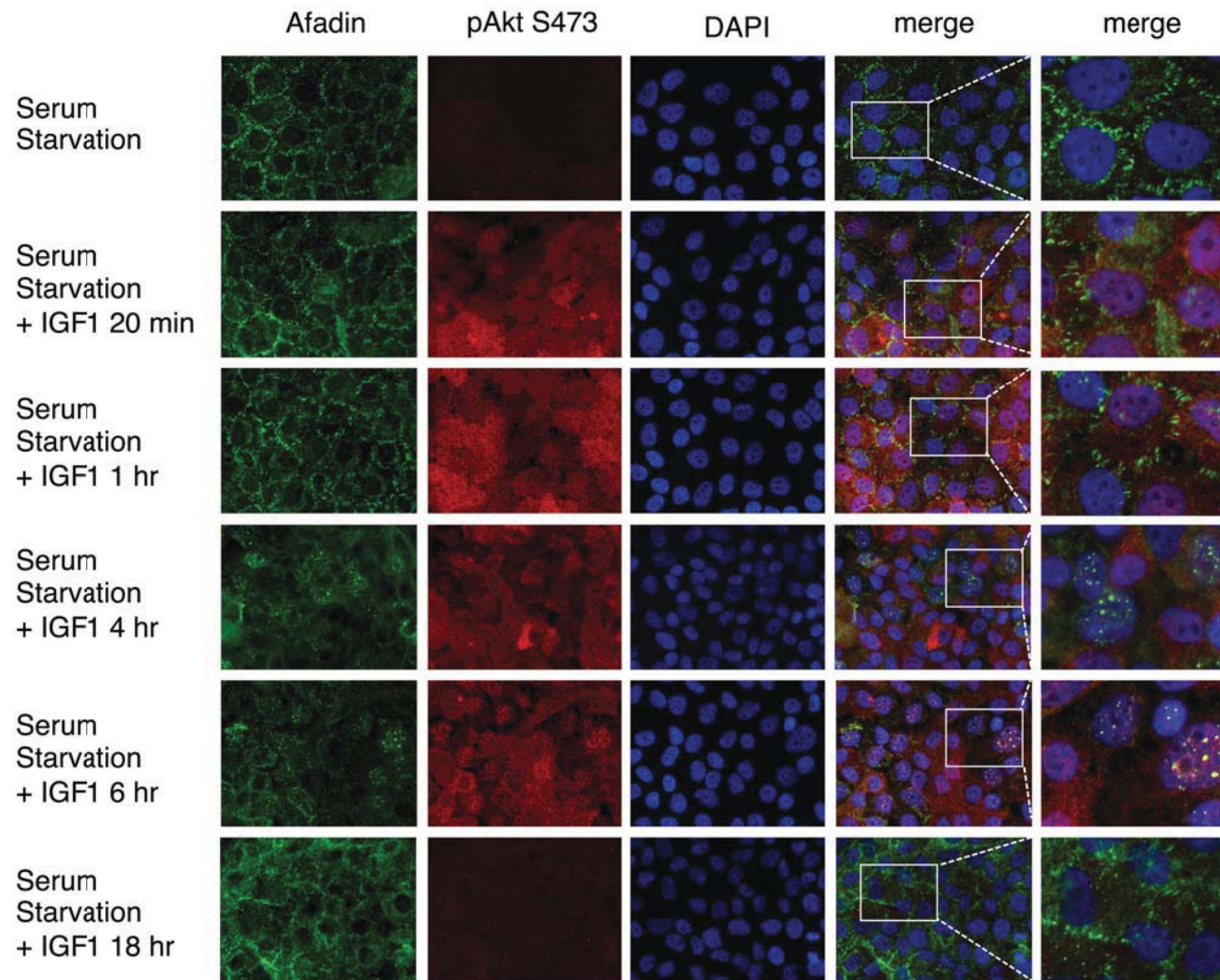
Fig. 7



Supplementary Fig. S1:

- A. Related to Fig. 1C: HeLa cells were serum starved, pre-treated with the indicated inhibitors and stimulated with IGF-1 for 20 min. In order to assess phosphorylation of Afadin at S1718, whole cell lysates were immunoblotted with the indicated antibodies.
- B. Related to Fig. 2A: Akt1 and Akt2 were silenced separately and concomitantly using shRNA in HeLa cells. Phosphorylation of Afadin was assessed. Whole cell lysates were immunoblotted with the indicated antibodies. HeLa cells do not express Akt3.
- C. Related to Fig. 2D: HeLa cells were transfected with pcDNA3, PIK3CA WT, PIK3CA H1047R, PIK3CA E545K. Cells were serum starved and phosphorylation of Afadin was determined using immunoblotting

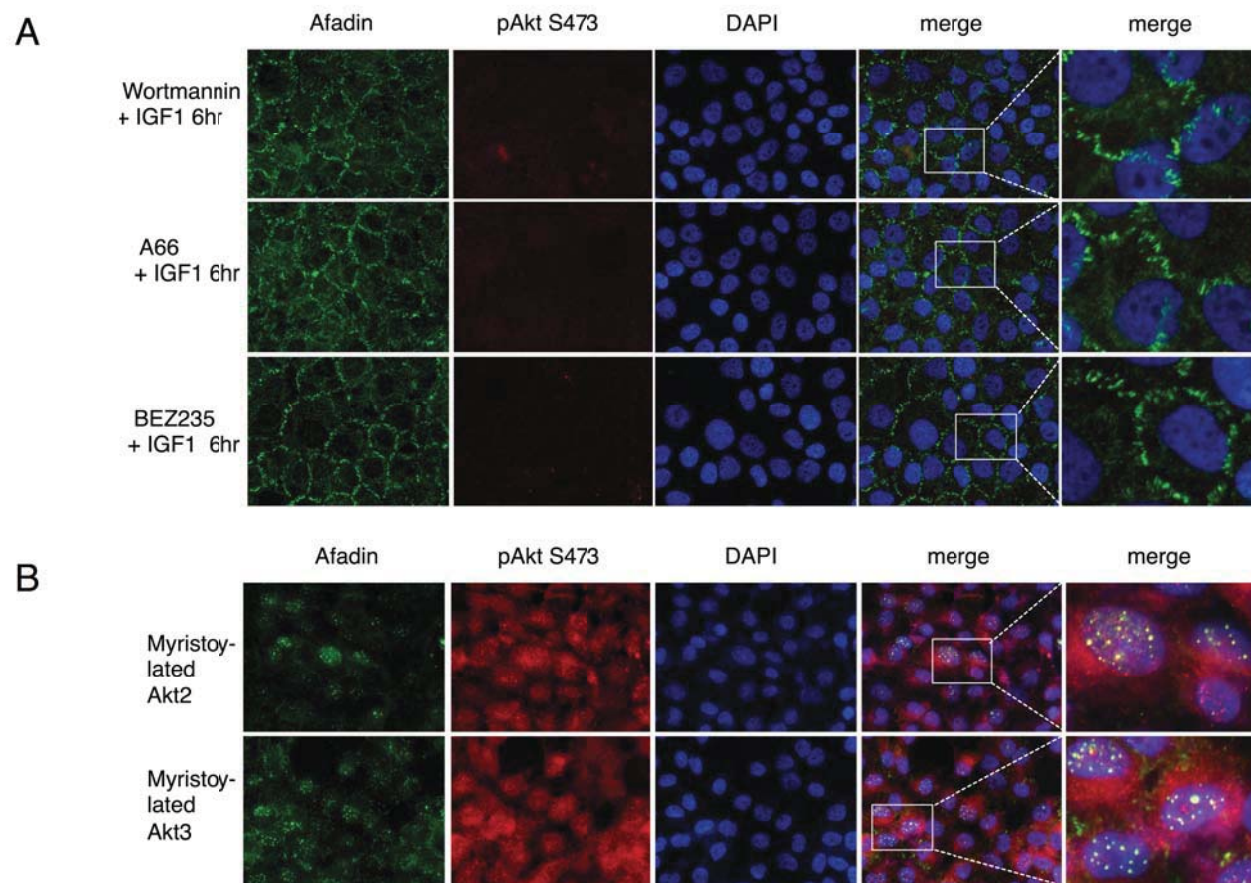
Supplementary Fig. S1



Supplementary Fig. S2:

Related to Fig. 3A: MCF10A cells were serum starved and stimulated with IGF-1 for 20 min, 1 hr, 4 hr, 6 hr and 18 hr. Intracellular localization of Afadin was assessed using immunofluorescence. Nuclei were stained with DAPI. Results are representative of multiple fields and at least three independent experiments

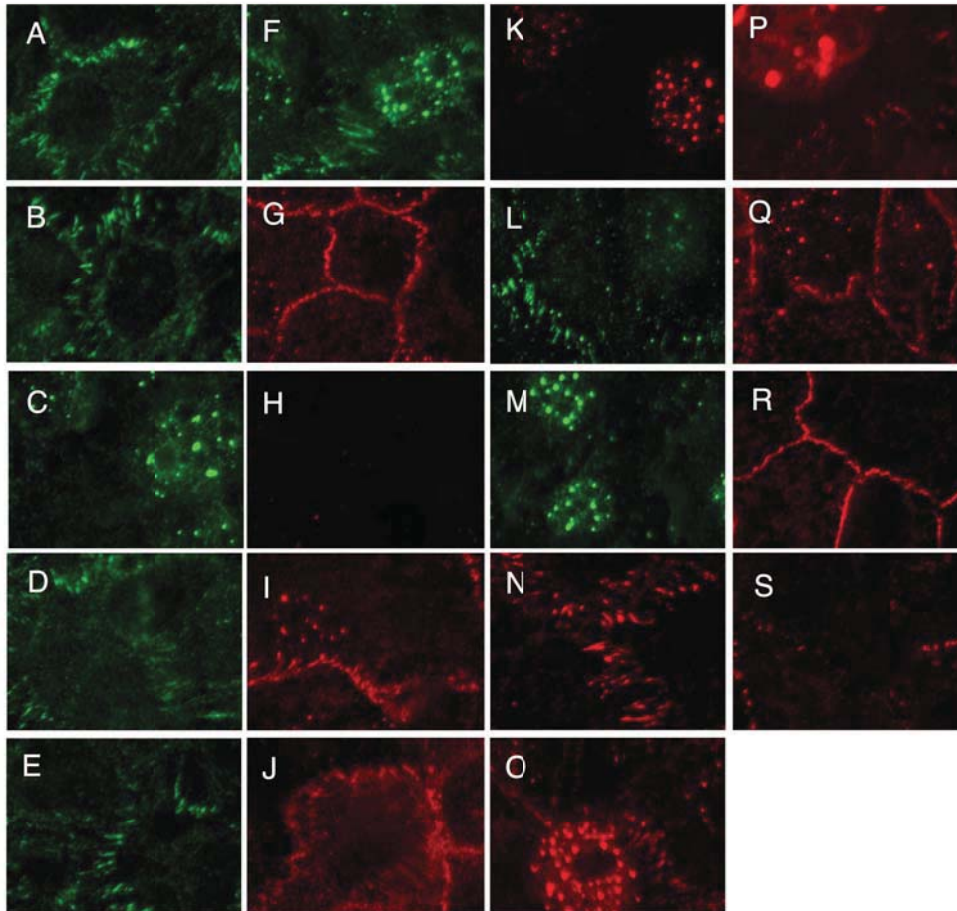
Supplementary Fig. S2



Supplementary Fig. S3:

- A. Related to Fig. 3A: MCF10A cells were serum starved and pretreated with the indicated inhibitors, and then stimulated with IGF-1 for 20 min. Intracellular localization of Afadin was assessed using immunofluorescence. Nuclei were stained with DAPI. Results are representative of multiple fields and at least three independent experiments.
- B. Related to Fig. 4B: MCF10A cells transfected with myristoylated Akt2 or Akt3 alleles and immunofluorescence performed with the indicated antibodies. In all cases nuclei were stained with DAPI. Images are representative of multiple fields, and of at least three independent experiments.

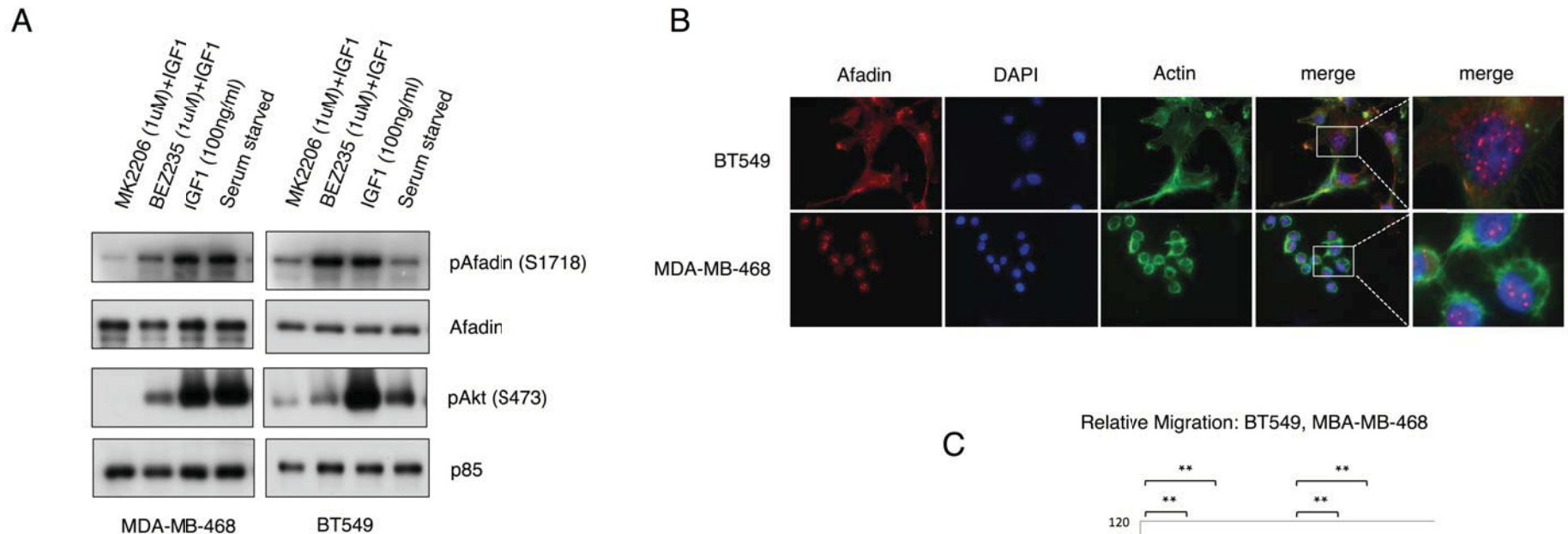
Supplementary Fig. S3



Supplementary Fig. S4:

Related to Fig. 3-5: Magnified single channel Afadin staining: **A-D** relate to Figure 3A: A: serum starvation; B: IGF1 1hr; C: IGF1 6 hr; D: MK2206 + IGF1 6 hr. **E-F** relate to Figure 3B: E: serum starvation; F: IGF1 6 hr. **G-K** relate to Figure 4A: G: pLKO; H: shAfadin; I: shAfadin + Afadin WT; J: shAfadin + Afadin S1718A; K: shAfadin + Afadin S1718E. **L-M**: relate to Figure 4B: L: pcDNA3; M: MyrAkt1. **N-S**: relate to Figure 5C: N: Serum starvation; O: IGF1 6 hr; P: IGF1 + CHX 6 hr; Q: Full serum; R: MK2206 6 hr; S: MK2206 + CHX 6 hr.

Supplementary Fig. S4

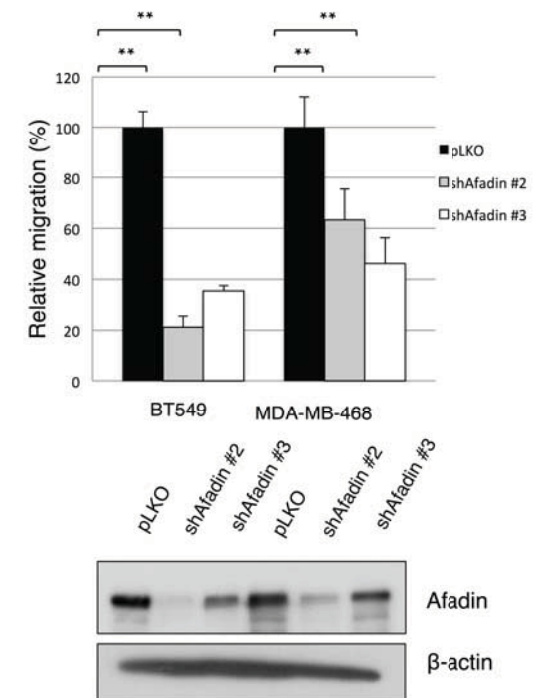


Supplementary Fig. S5, Related to Fig. 6:

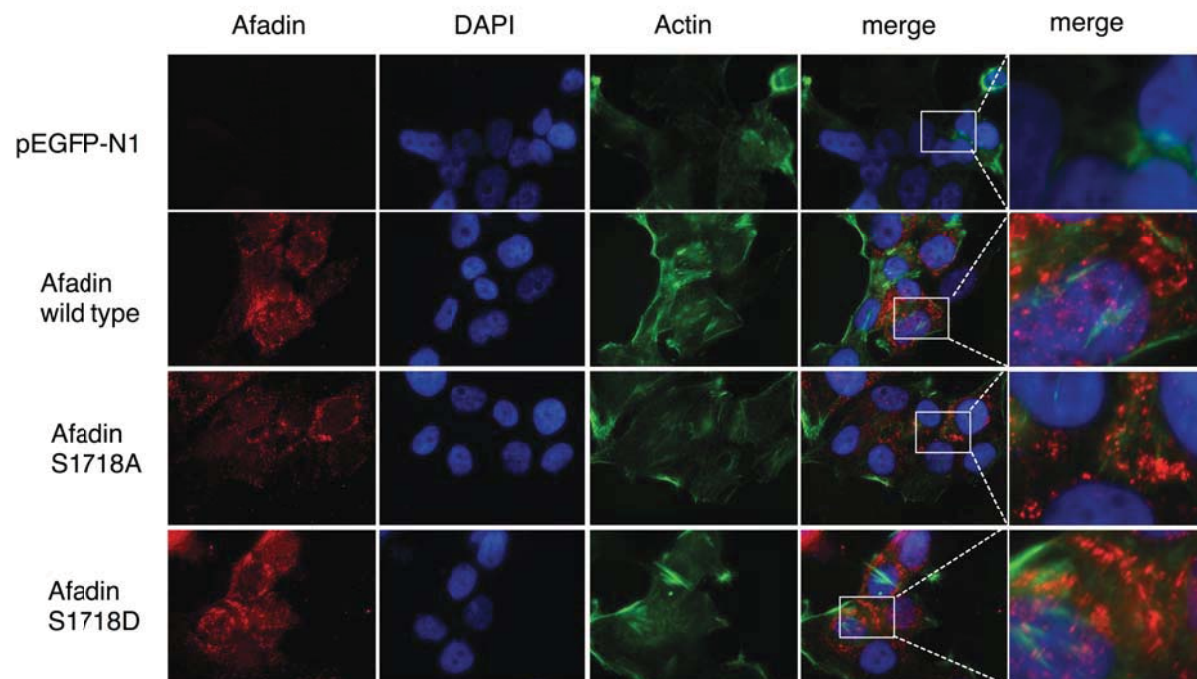
- A. BT549 and MDA-MB-468 cells were serum starved for 18 hr then stimulated with IGF-1 for 20 min either alone or after treatment with BEZ235 and MK2206 for 20 min. Phosphorylation of Afadin was evaluated by immunoblotting.
- B. BT549 and MDA-MB-468 cells were serum starved for 18 hr. Intracellular localization of Afadin was assessed using immunofluorescence. Results are representatives of multiple fields and at least three independent experiments.
- C. BT549 and MDA-MB-468 cells were infected with shRNA to silence Afadin (two different shRNA sequences; shAfadin #2 and shAfadin #3). Transwell cell migration was assessed. Expression of Afadin following silencing was measured using immunoblotting. ** $p < 0.01$ (Student T-test).

C

Relative Migration: BT549, MDA-MB-468



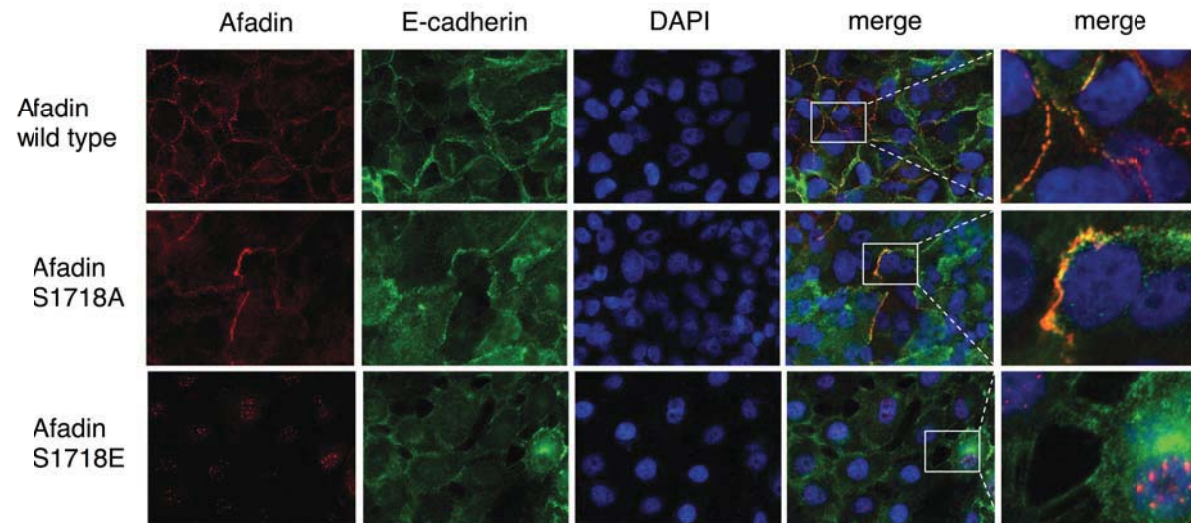
Supplementary Fig. S5



Supplementary Fig. S6 related to Fig. 6:

T47D transfected with pEGFP-N1, wild-type (WT) Afadin, Ser1718Ala (S1718A), Ser1718Asp (S1718D) Afadin. Immunofluorescence staining was performed using the indicated antibodies. Nuclei were stained with DAPI. Results are representative of multiple fields and at least three independent experiments

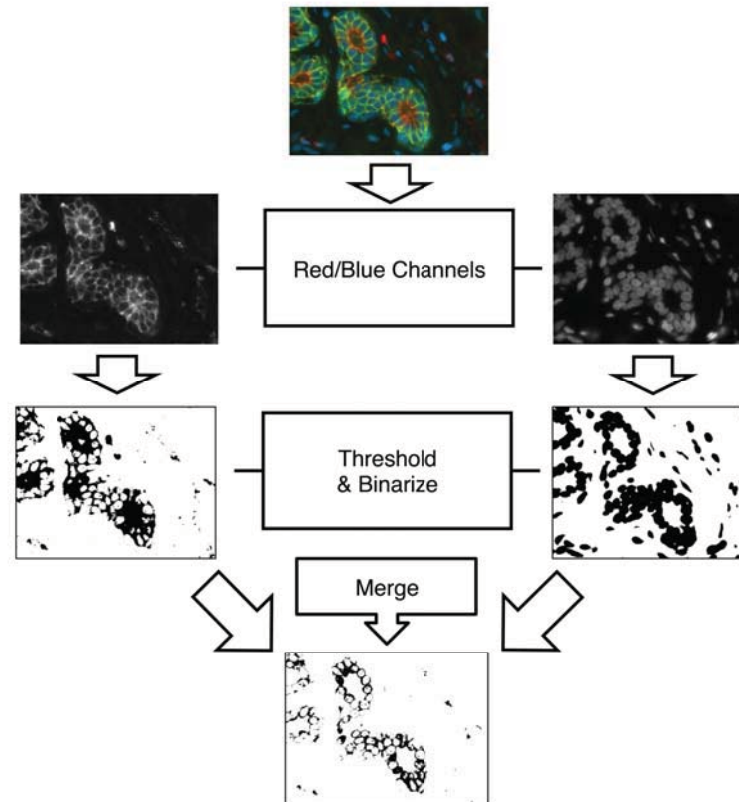
Supplementary Fig. S6



Supplementary Fig. S7 related to Fig. 6C:

MCF10A cells were transduced with shRNA to silence Afadin and transfected with wild type Afadin wild type, Ser1718Ala Afadin (S1718A) or Ser1718Glu Afadin (S1718E). The cells were evaluated by immunofluorescence with the indicated antibodies. Results are representative of multiple fields and at least three independent experiments

Supplementary Fig. S7



Supplementary Fig. S8 related to Fig. 7

Schematic representation of the quantification of Afadin nuclear localization. To quantitatively assess nuclear localization of Afadin, the immunofluorescence images were split into red, green, and blue channels and background subtraction was performed on the red and blue channels using a sliding paraboloid with a 50 pixel radius. After background correction, we applied an auto-threshold function to create binary images from the red and blue channels. The red channel represents Afadin staining, and the blue channel represents nuclear staining. We computed the total area of red and blue as well as the area of overlap of the red and blue binary images. We computed the Afadin nuclear localization score as the (red and blue overlapping area)/(total red area + total blue area).

Supplementary Fig. S8